Cellular Expression of $\beta_2 AR - \beta gal \Delta \alpha$ Fusion Protein in C2 Clones (measured by anti- β -gal ELISA)

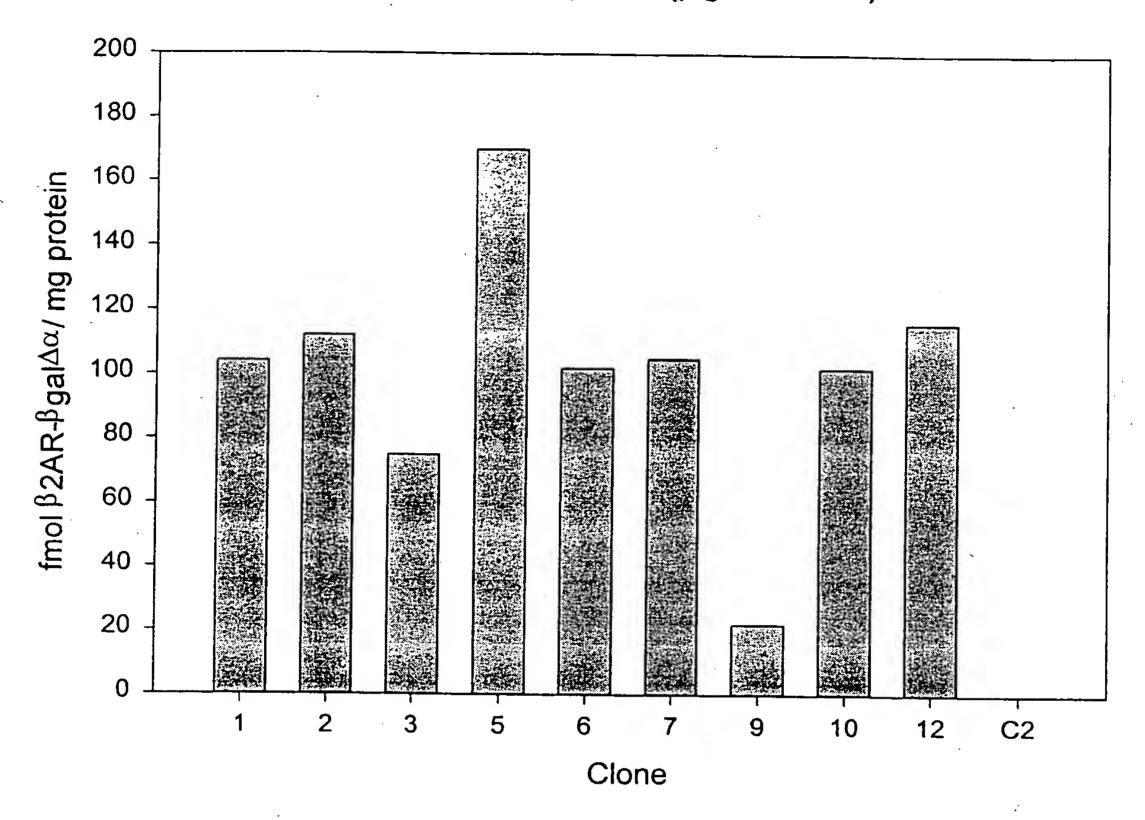


FIGURE 1A

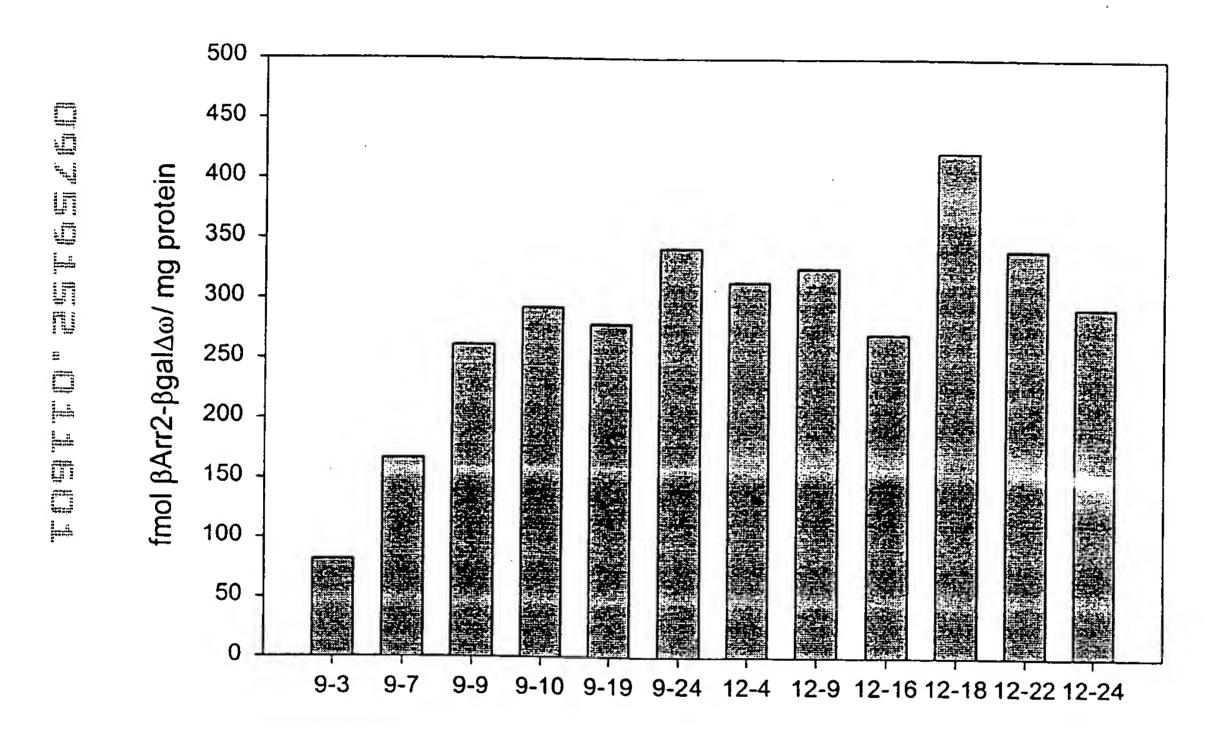


FIGURE 1B

Agonist Stimulated cAMP Response in C2 Cells Expressing $\beta 2AR\text{-}\beta gal\Delta\alpha$

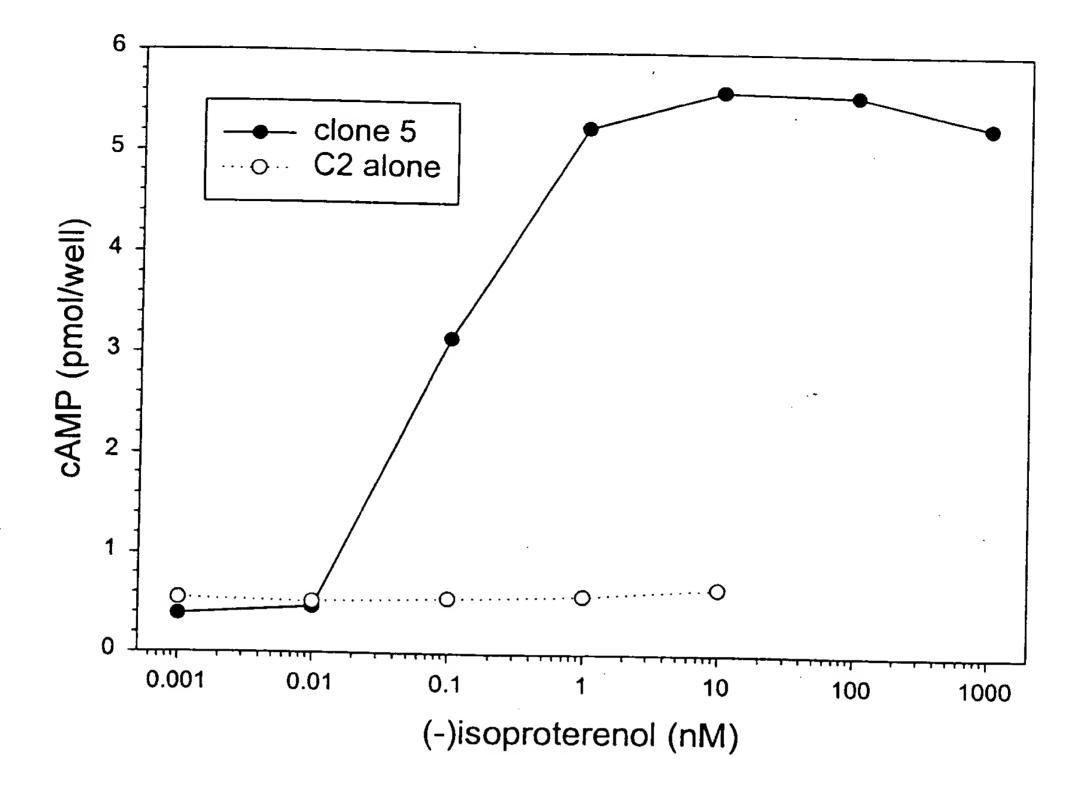
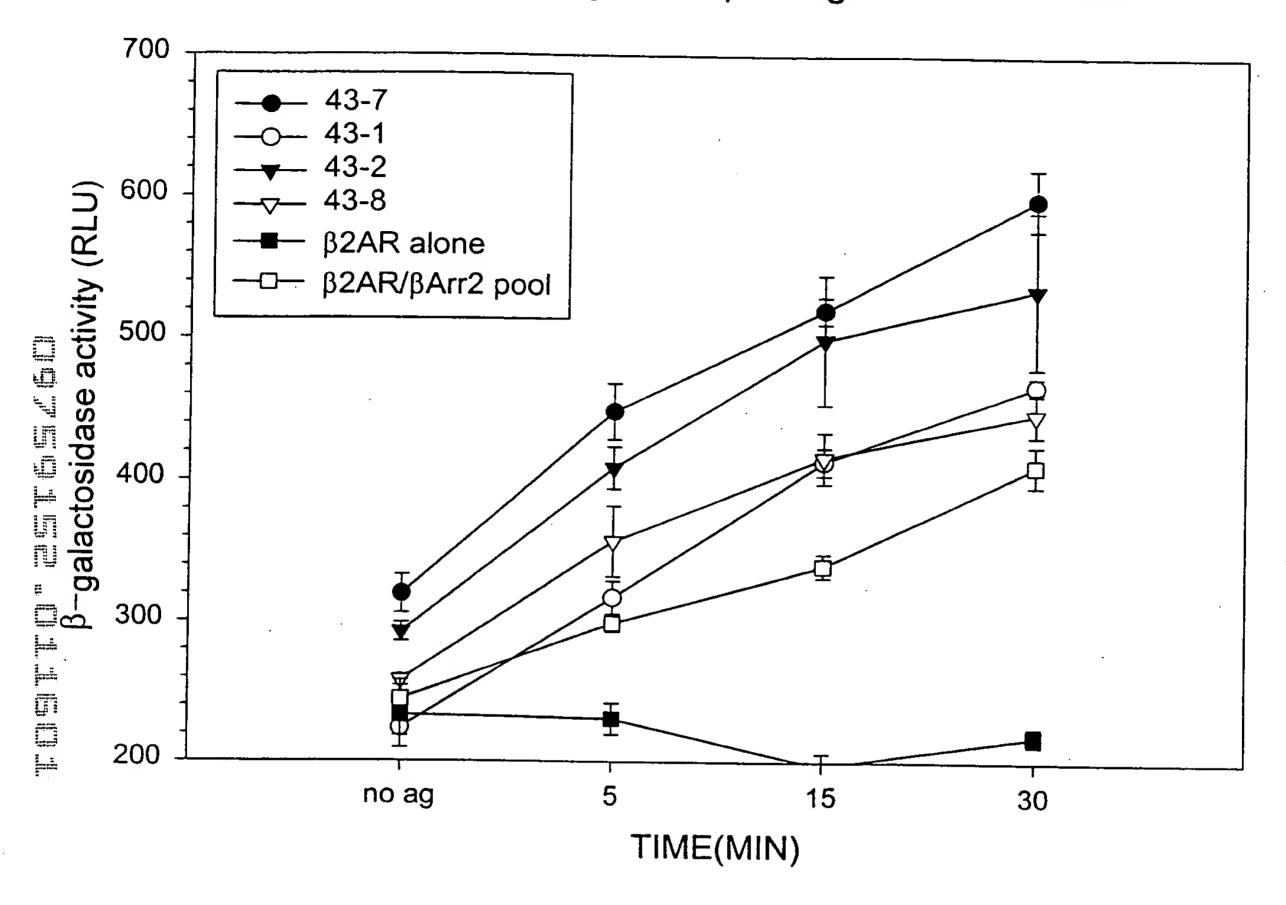


FIGURE 2

$\beta-$ galactosidase Complementation as a Measurement for $\beta 2AR-\beta gal\Delta\alpha$ interacting with $\beta Arrestin 2-\beta gal\Delta\omega$ upon agonist Stimulation



 β –galactosidase Complementation as a Measurement for $\beta 2AR$ - $\beta gal\Delta\alpha$ Interaction with $\beta Arrestin 1$ - $\beta gal\Delta\omega$ upon Agonist Stimulation

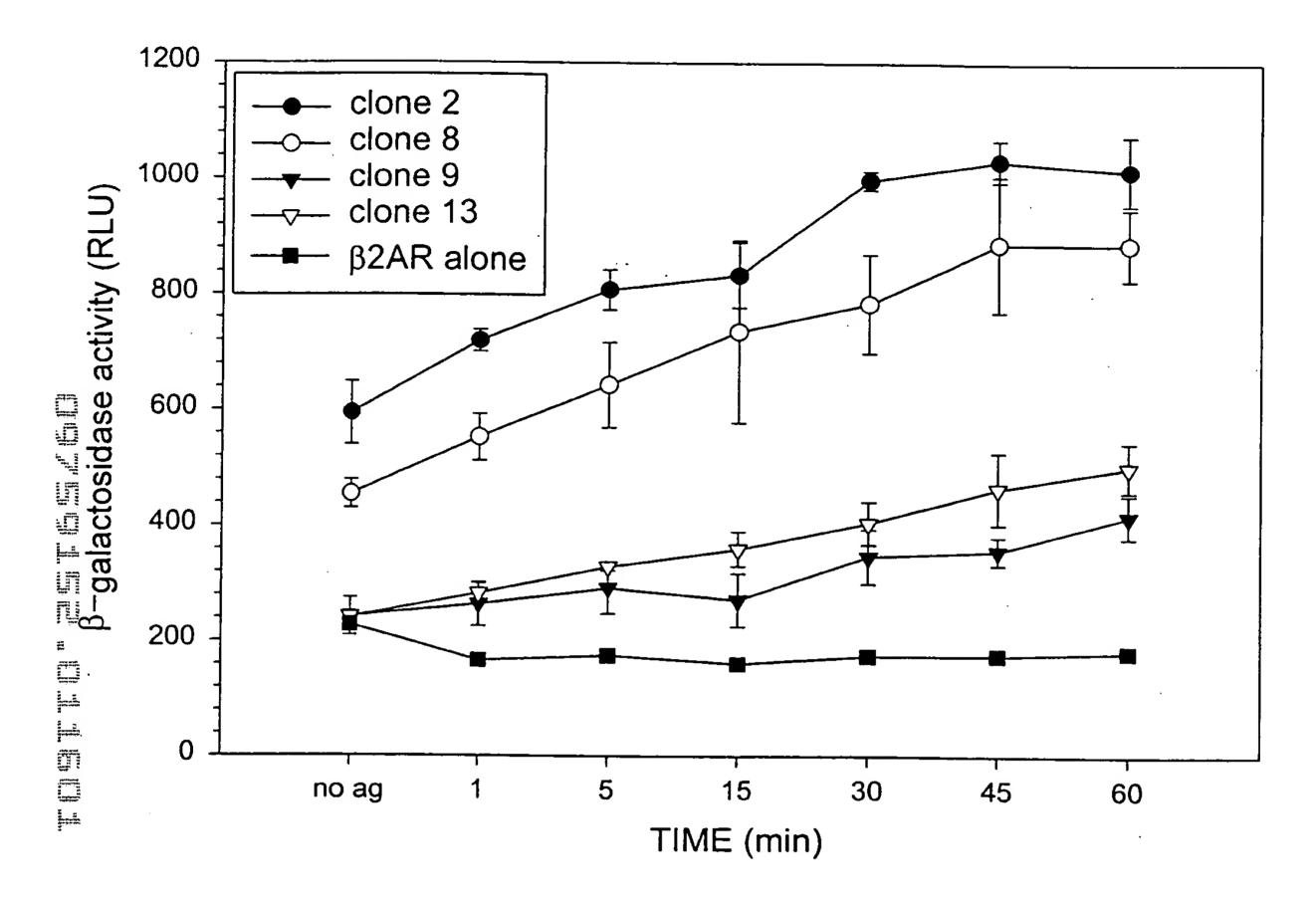


FIGURE 3B

β –galactosidase Activity in Response to Agonist in C2 Cells Coexpressing $\beta 2AR$ - $\beta gal\Delta\alpha$ and $\beta Arrestin 2$ - $\beta gal\Delta\omega$ Fusion Proteins

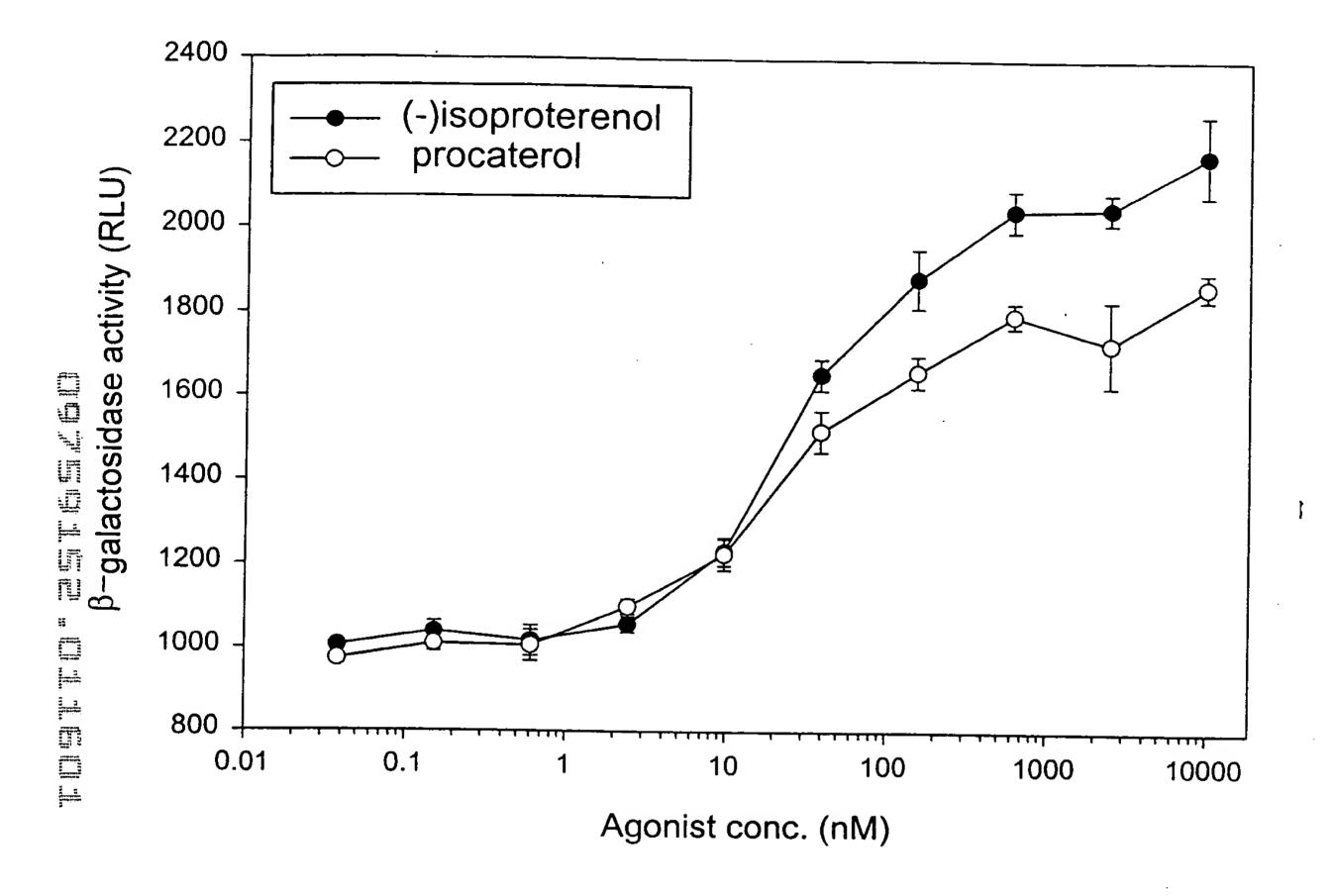


FIGURE 4A

β –galactosidase Activity in Response to Agonist in C2 Cells Coexpressing $\beta 2AR$ - $\beta gal\Delta\alpha$ and $\beta Arrestin1$ - $\beta gal\Delta\omega$ Fusion Proteins

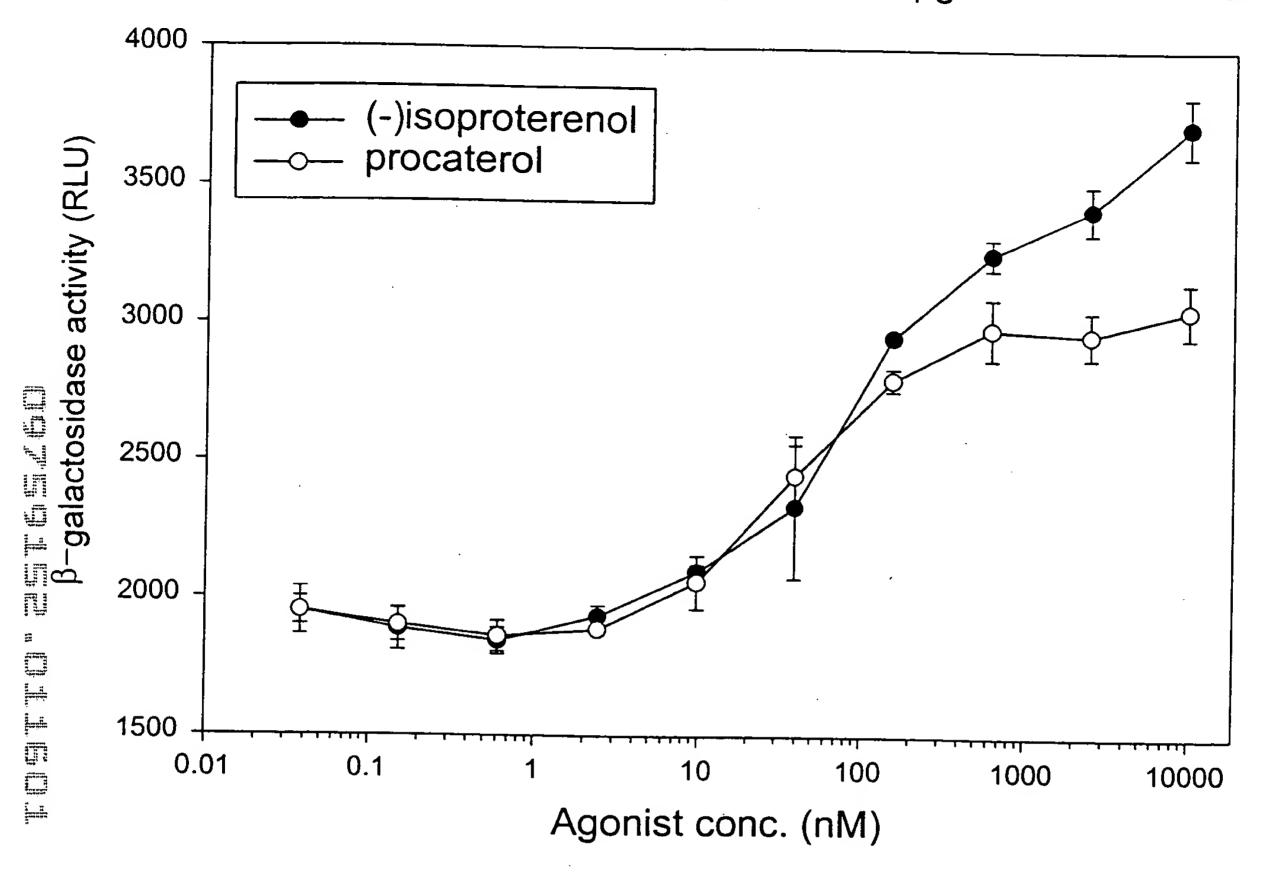
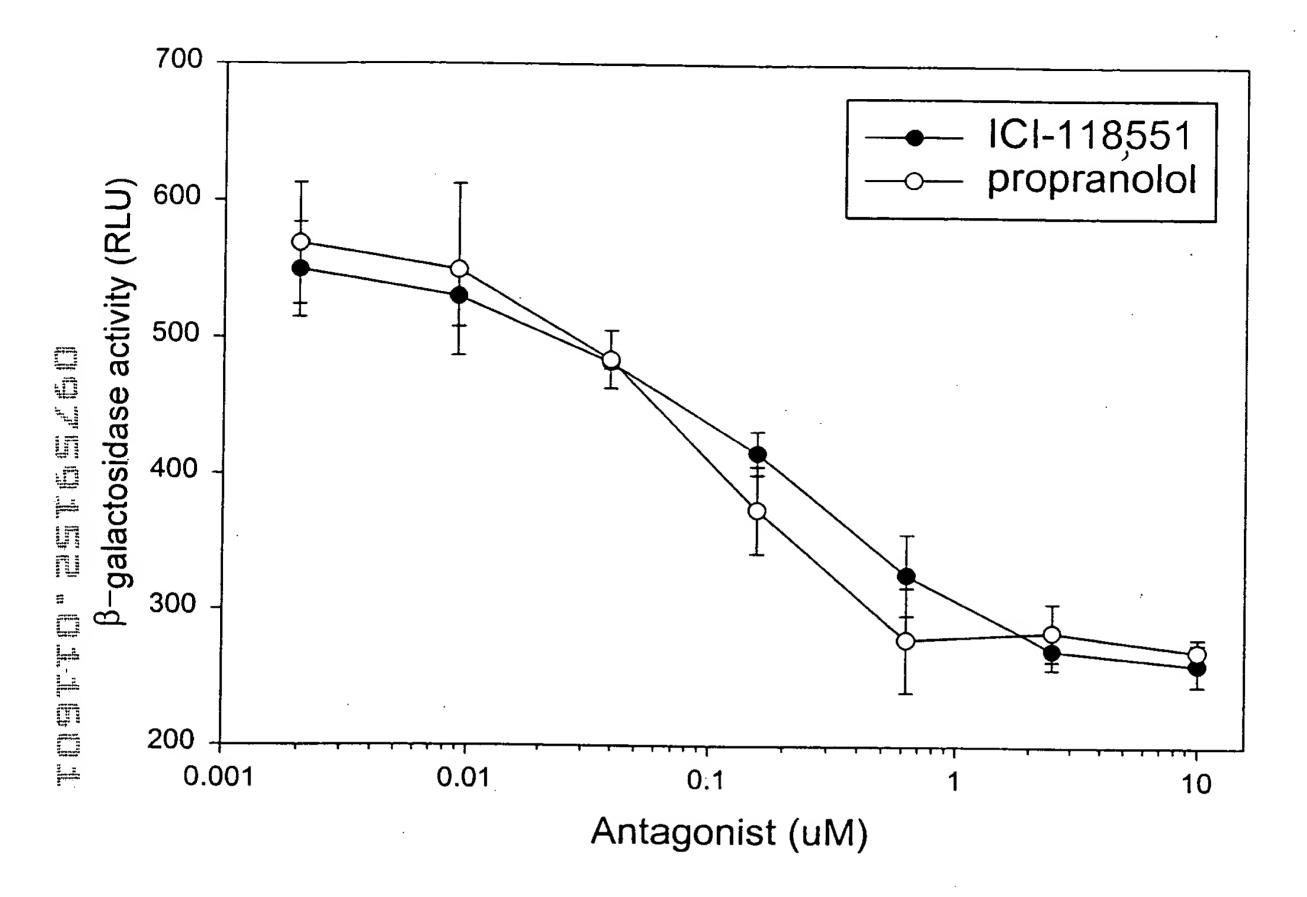


FIGURE 4B

Inhibition of β –galactosidase activity in C2 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arrestin 2-\beta gal\Delta\omega$ Fusion Proteins



Antagonist Inhibition of β -galactosidase Activity in C2 Cells Coexpressing β 2AR- β gal $\Delta\alpha$ and β Arrestin1- β gal $\Delta\omega$ Fusion Proteins

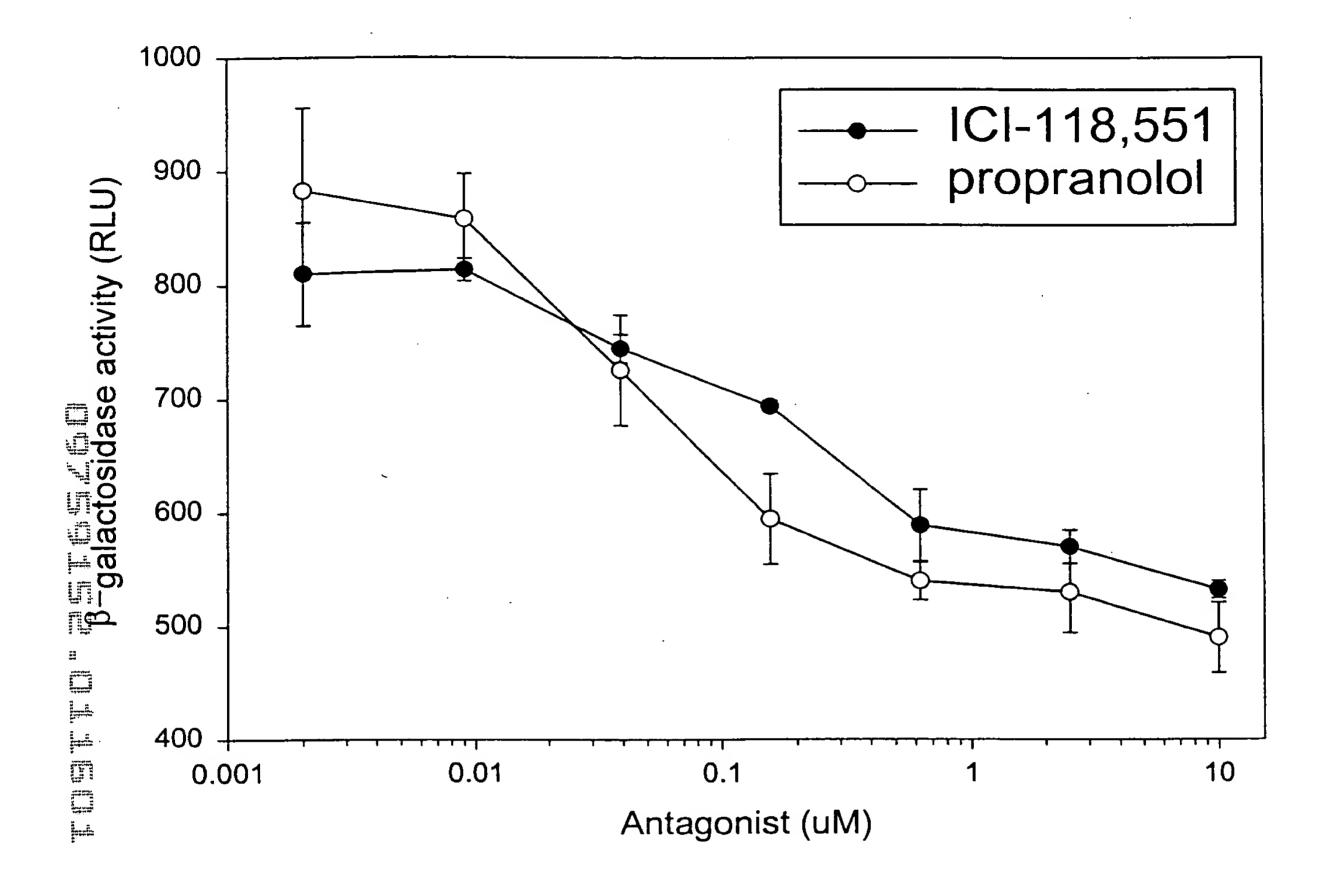


Figure 5B

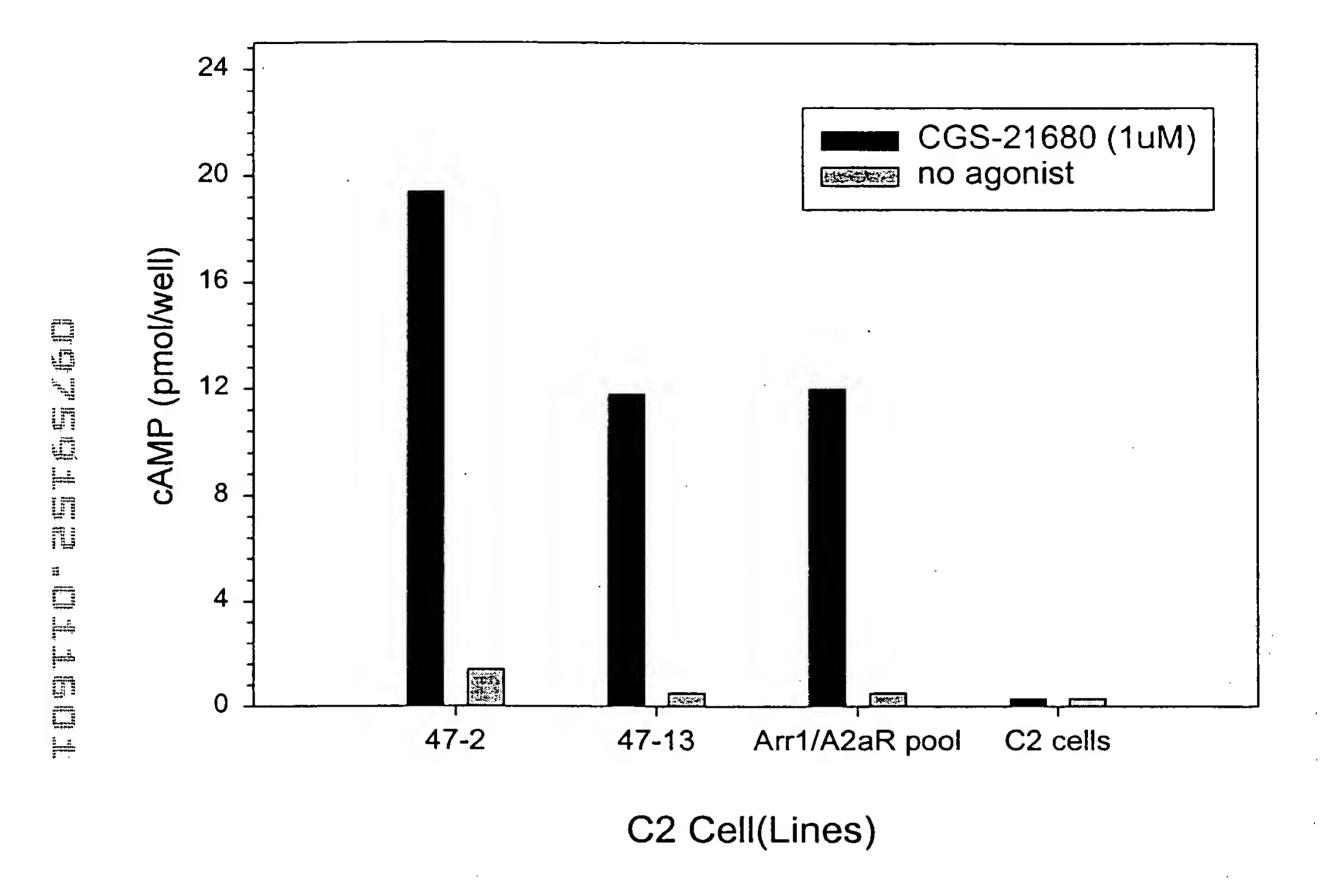


FIGURE 6

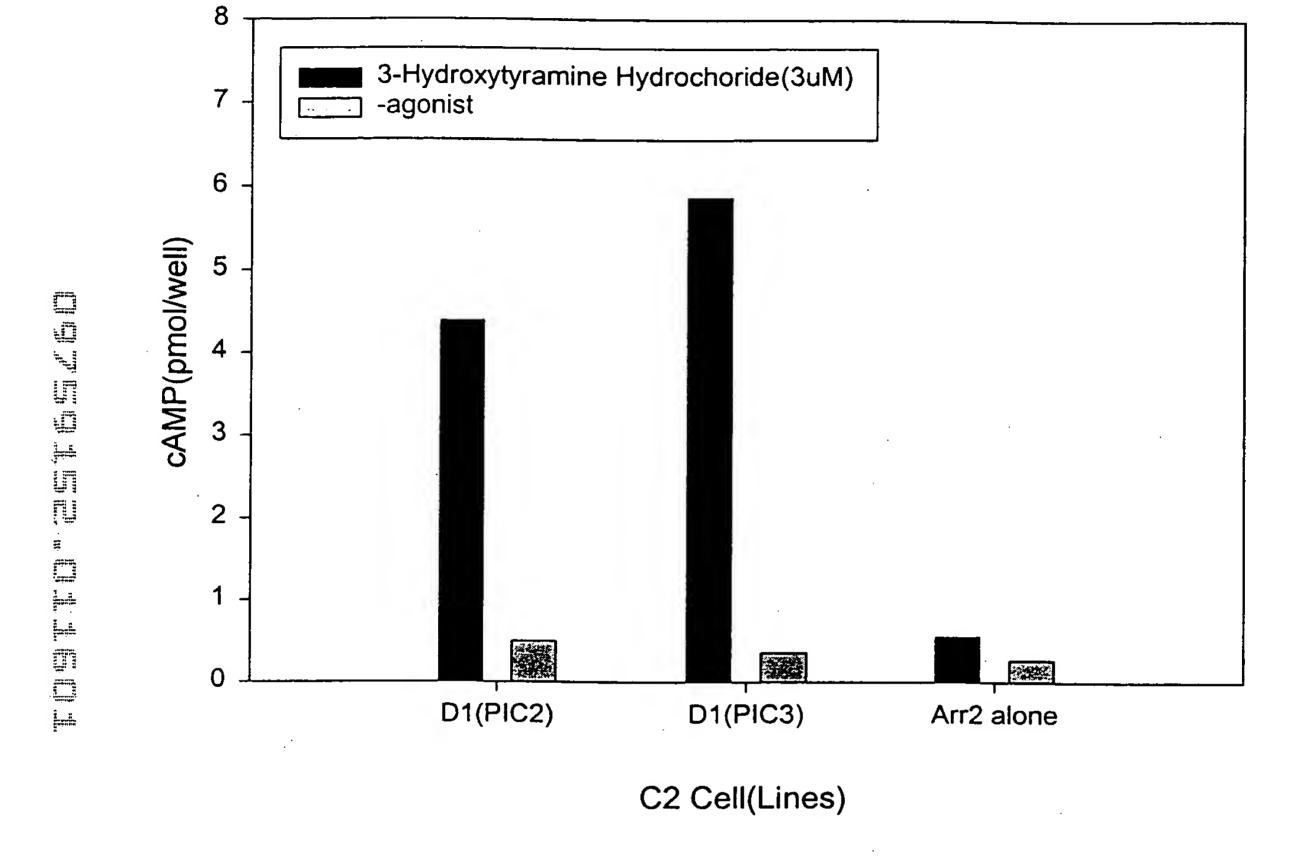


FIGURE 7

$β_2AR-βgalΔω$ and βarr2-βgalΔα Interaction in HEK293 Clones in Response to Isoproterenol Treatment (1 μM)

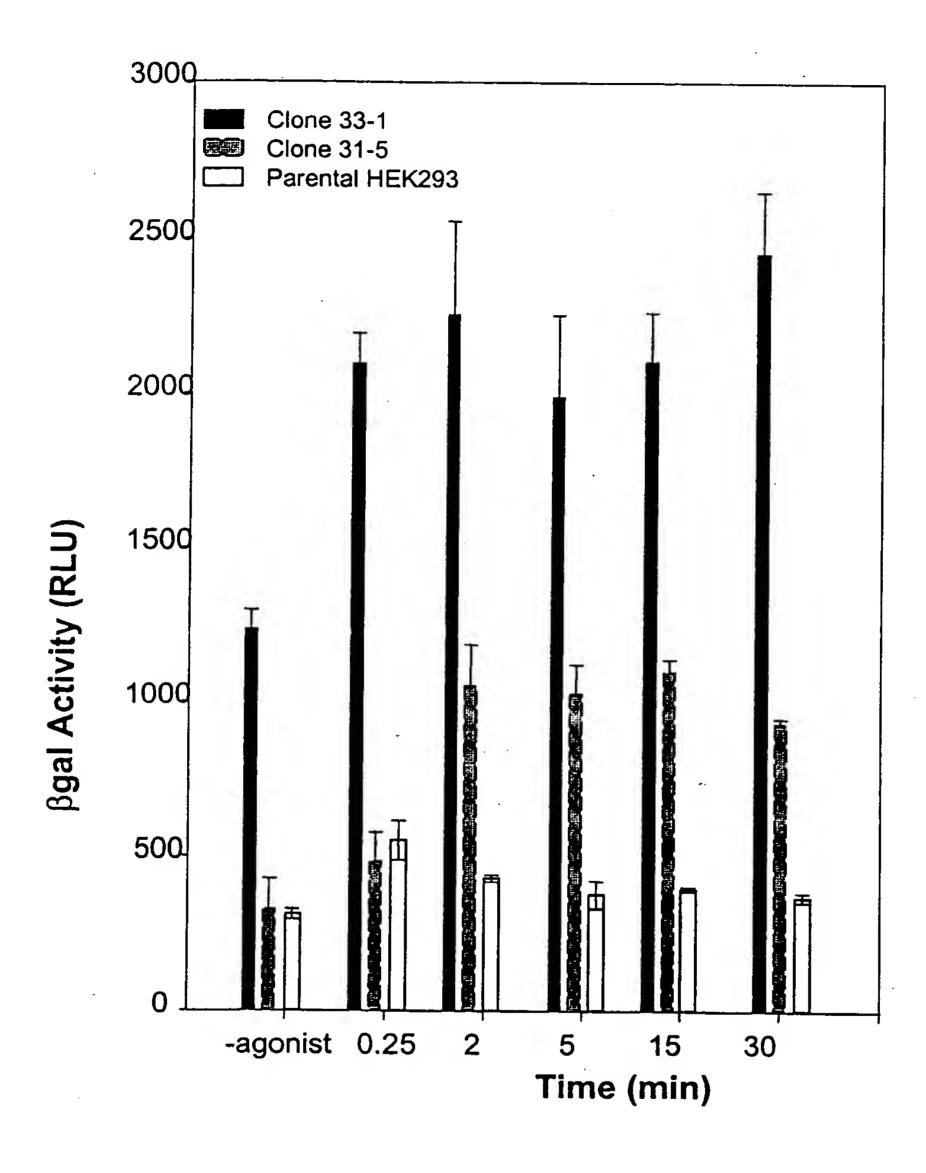
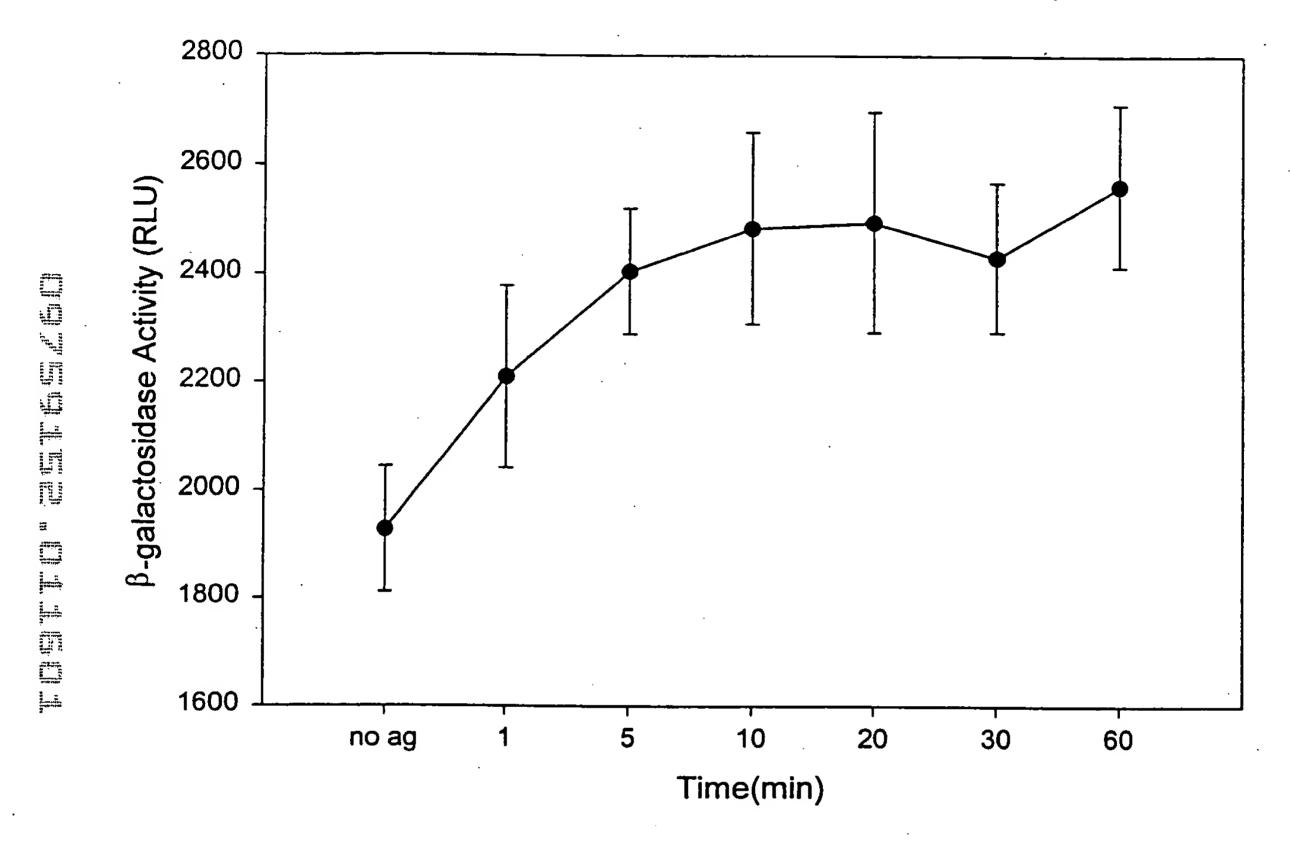
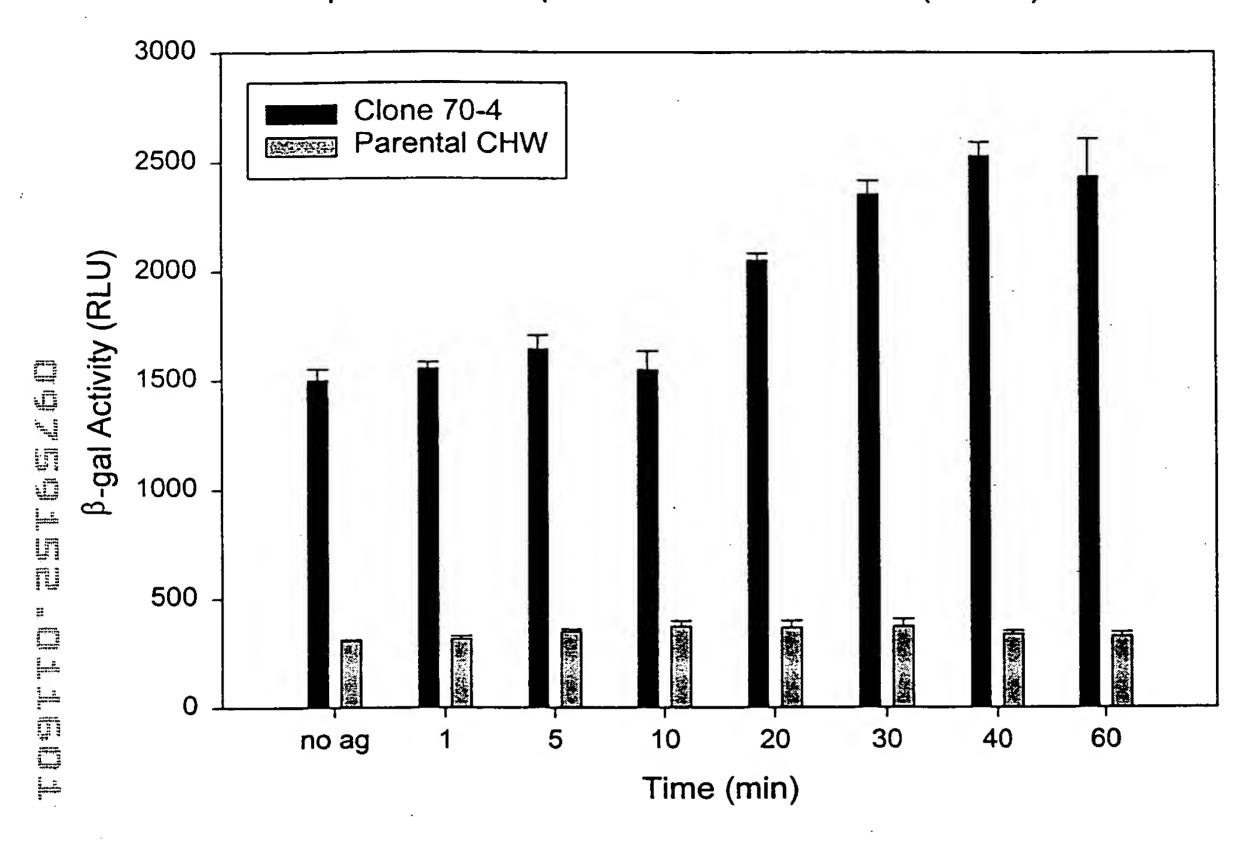


FIGURE 8A

 $\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arr1-\beta gal\Delta$ ω Interaction in a CHO Pool in Response to Isoproterenol Treatment(10uM)



$\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arr2-\beta gal\Delta\omega$ Interaction in CHW Clone in Response to Isoproterenol Treatment (10uM)



 β –galactosidase Complementation as a Measurement for Adrenergic Receptor Homodimerization in HEK 293 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta 2AR-\beta gal\Delta\omega$.

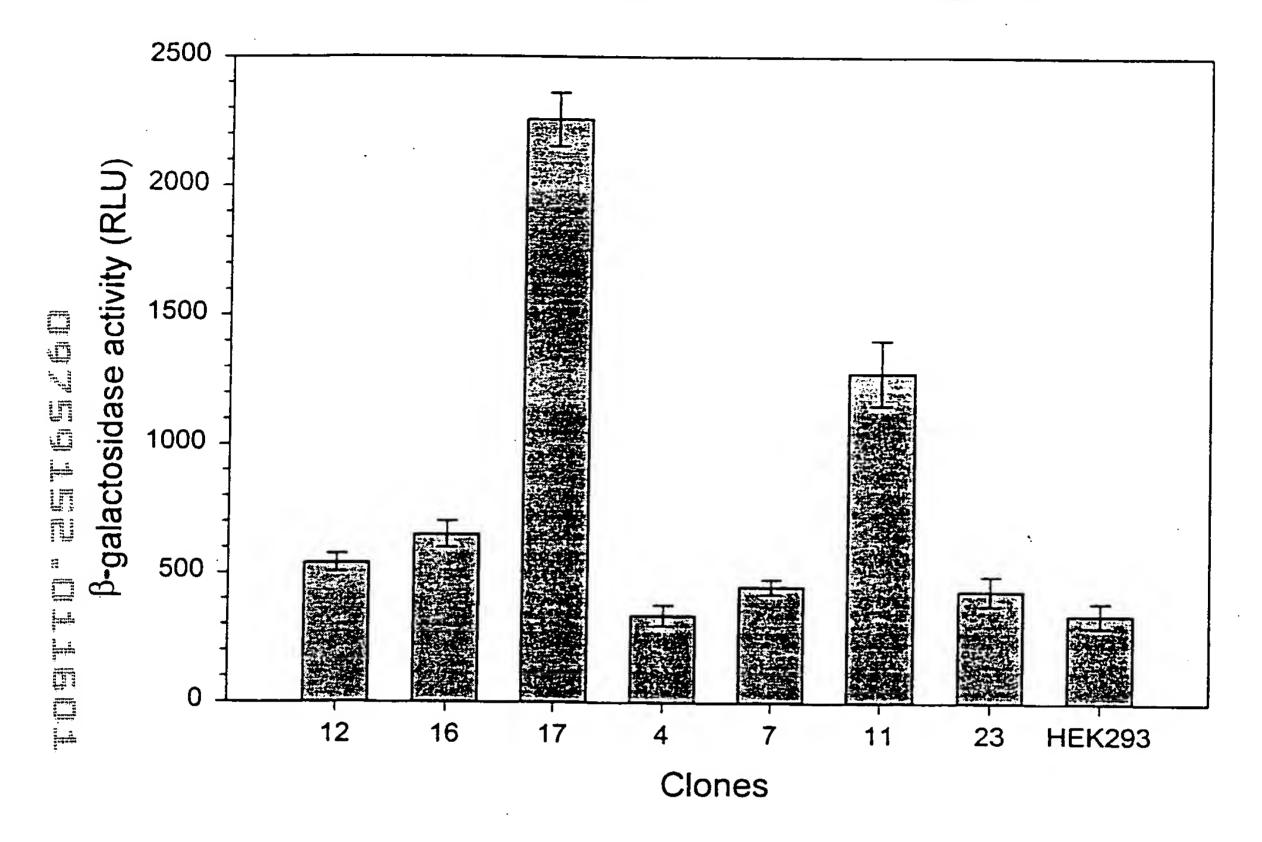
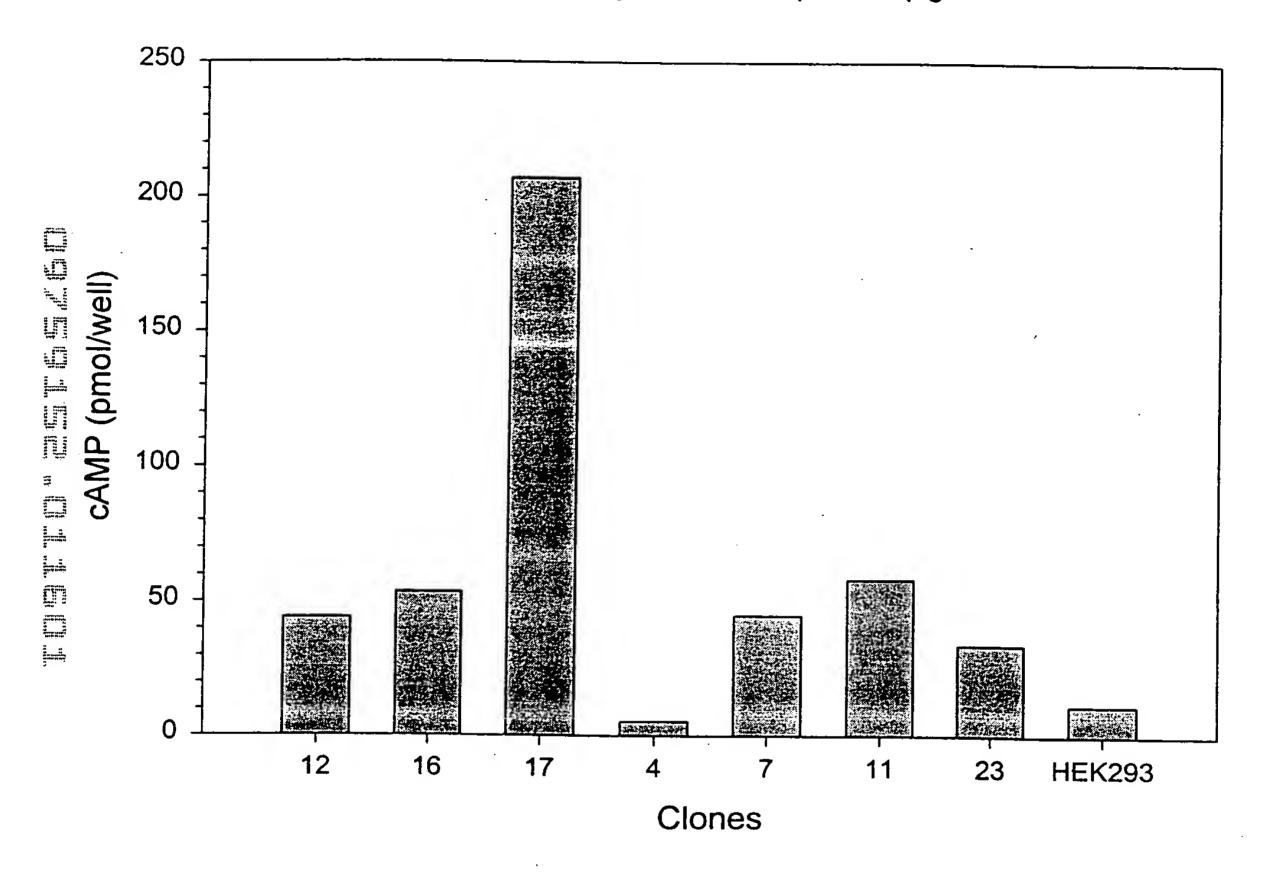


FIGURE 9A

Agonist Stimulated cAMP Response in HEK 293 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta 2AR-\beta gal\Delta\omega$



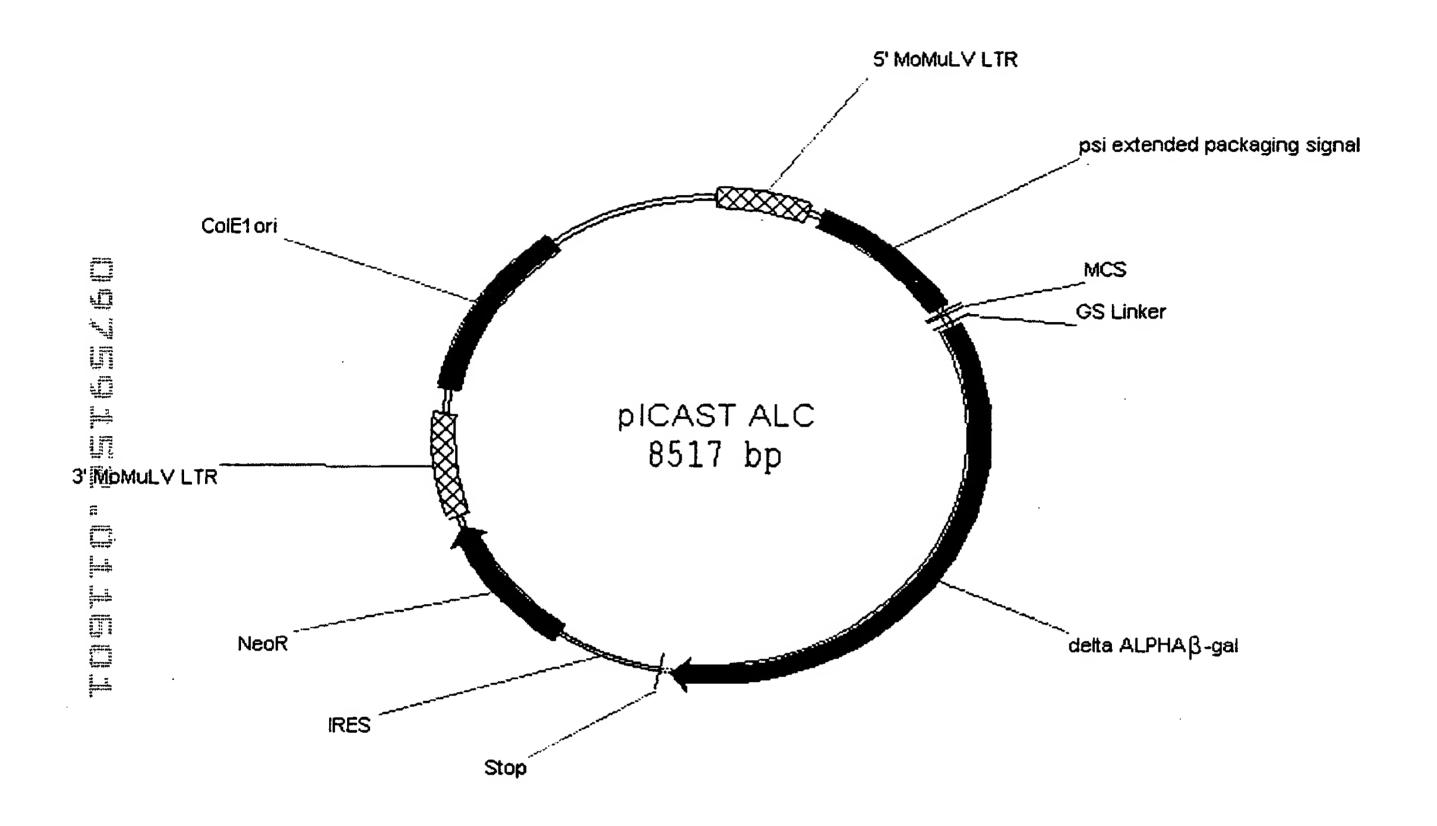


Figure 10A

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101	GGATATCTGT CCTATAGACA					
151	GGTCCCCAGA CCAGGGGTCT					
201	GTTTCCAGGG CAAAGGTCCC					
251	TAAÇCAATCA ATTGGTTAGT					
301 1	GCTCAATAAA CGAGTTATTT					- -
351 4	ACTGACTCAG	CGGGCCCATG	GGCACATAGG	TTATTTGGGA	CTTGCAGTTG GAACGTCAAC	
401 []	GTAGGCTGAA	CACCAGAGCG	ACAAGGAACC	CTCCCAGAGG	TCTGAGTGAT AGACTCACTA	
15 1			AGAAAGTAAA	CCCCGAGCA	CCGGGATCGG GGCCCTAGCC	
591 <u>5</u> 91	GAGACCCCTG CTCTGGGGAC				CAAGCTGGCC GTTCGACCGG	
551	AGCAACTTAT TCGTTGAATA		GGCTAACAGA	TCACAGATAC	TGACTAAAAT	
601	TGCGCCTGCG ACGCGGACGC				CTGGCGGACC GACCGCCTGG	-
651	CGTGGTGGAA GCACCACCTT				CTGGGAGACG GACCCTCTGC	
701	TCCCAGGGAC AGGGTCCCTG				GGAAGGGAGT CCTTCCCTCA	
751	_ 	AGGCTGGGGC	AGTCCTATAC	ACCAAGACCA	TCCTCTGCTC	
801		TCAAGGGCGG	AGGCAGACTT	AAAAACGAAA	GCCAAACCTT	
851	CCGAAGCCGC GGCTTCGGCG	GCGTCTTGTC CGCAGAACAG	TGCTGCAGCA ACGACGTCGT	TCGTTCTGTG AGCAAGACAC	TTGTCTCTGT AACAGAGACA	
901	CTGACTGTGT	TTCTGTATTT	GTCTGAAAAT	TAGGGCCAGA		

951				AGATGTCGAG TCTACAGCTC		
1001	· ·			GTTGGGTTAC CAACCCAATG		
1051				CCGCGAGACG GGCGCTCTGC		
1101				GGTCTTTTCA CCAGAAAAGT		
1151				TGACÇTGGGA ACTGGACCCT		
1201		GAGGGACCCA	GTTCGGGAAA	GTACACCCTA CATGTGGGAT		
1251 11 12				CCTTGAACCT GGAACTTGGA		
13701 40 144				TCACTCCTTC AGTGAGGAAG		
1 35 1				ATAGGGCGAT TATCCCGCTA		
				AATTGGGAGG TTAACCCTCC		
+2 1 45 1] CTCGAGATGG	GCGTGATTAC	GGATTCACTG	A V V A	CCCGCACCGA	
+2				G E W		
1501				TGGCGAATGG ACCGCTTACC		·
+2	W F P A	P E A	V P E	S W L E	C D L	
1551				GCTGGCTGGA CGACCGACCT		
+2	PEAI	T V V	V P S	и у W и	M H G Y	
1601				AACTGGCAGA TTGACCGTCT		
+2	D A P	I Y T	V T Y	PIT	V N P	
1651				TCCCATTACG AGGGTAATGC		· · · · · · · · · · · · · · · · · · ·
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+2 P F V P T E N P T G C Y S L T F N
1701 CGTTTGTTCC CACGGAGAAT CCGACGGGTT GTTACTCGCT CACATTTAAT GCAAACAAGG GTGCCTCTTA GGCTGCCCAA CAATGAGCGA GTGTAAATTA
+2 V D E S W L Q E G Q T R I I F D G
1751 GTTGATGAAA GCTGGCTACA GGAAGGCCAG ACGCGAATTA TTTTTGATGG CAACTACTTT CGACCGATGT CCTTCCGGTC TGCGCTTAAT AAAAACTACC
+2 V N S A F H L W C N G R W V G Y
1801 CGTTAACTCG GCGTTTCATC TGTGGTGCAA CGGGCGCTGG GTCGGTTACG GCAATTGAGC CGCAAAGTAG ACACCACGTT GCCCGCGACC CAGCCAATGC
+2GQDSRLPSEFDLSAFLR
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1901 GCCGGAGAAA ACCGCCTCGC GGTGATGGTG CTGCGCTGGA GTGACGGCAG CGGCCTCTTT TGGCGGAGCG CCACTACCAC GACGCGACCT CACTGCCGTC
T+2 Y L E D Q D M W R M S G I F R D
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+2 V S L L H K P T T Q I S D F H V A
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+2 M C G E L R D Y L R V T V S L W
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+2 Q G E T Q V A S G T A P F G G E I
2151 AGGGTGAAAC GCAGGTCGCC AGCGGCACCG CGCCTTTCGG CGGTGAAATT TCCCACTTTG CGTCCAGCGG TCGCCGTGGC GCGGAAAGCC GCCACTTTAA
+2 I D E R G G Y A D R V T L R L N V
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+2 E N P K L W S A E I P N L Y R A
2251 CGAAAACCCG AAACTGTGGA GCGCCGAAAT CCCGAATCTC TATCGTGCGG GCTTTTGGGC TTTGACACCT CGCGGCTTTA GGGCTTAGAG ATAGCACGCC

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	ACCAACTTGA CGTGTGGCGG CTGCCGTGCG ACTAACTTCG TCTTCGGACG	
	D V C D D D V D	·
72	DVGFREVRIENGLLLLN	
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	CTACAGCCAA AGGCGCTCCA CGCCTAACTT TTACCAGACG ACGACGACTT	
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2401	CGGCAAGCCG TTGCTGATTC GAGGCGTTAA CCGTCACGAG CATCATCCTC GCCGTTCGGC AACGACTAAG CTCCGCAATT GGCAGTGCTC GTAGTAGGAG	
+2	L H G Q V M D E Q T M V Q D I L L	
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To the	ACGTACCAGT CCAGTACCTA CTCGTCTGCT ACCACGTCCT ATAGGACGAC	
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	TACTTCGTCT TGTTGAAATT GCGGCACGCG ACAAGCGTAA TAGGCTTGGT	
F + 2	PLWYTLC DRYGLYVD	
2551		
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+== 	~	
* +2	DPRWLPAMSERVTRMVQ	
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+2	L G G	F A K	 Y W O A	F R O	 У Р R	
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+2	L Q G G	F V W	D W V	 D Q S I	I K Y	
3051	TACAGGGCGG ATGTCCCGCC	CTTCGTCTGG GAAGCAGACC	GACTGGGTGG CTGACCCACC	ATCAGTCGCT TAGTCAGCGA	GATTAAATAT CTAATTTATA	
+2	DEN	G N P W	S A Y	G G D	F G D T	
3 10 1 卓					TTGGCGATAC AACCGCTATG	
## ##-2	P N D	R Q F C	M N G	L V F	A D R	
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+2	трнр	A L T	EAK	i QQQ	F F Q	
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+2	FRLS	G Q T	I E V	T S E	Y L F R	
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+2	P L A S	G E V	P L D V	A P Q	G K Q	
351	CGCTGGCAAG GCGACCGTTC	CGGTGAAGTG	 CCTCTGGATG	TCGCTCCACA	AGGTAAACAG	
+2	L I E L	P E L	P Q P	E S A (G Q L W	
401	TTGATTGAAC AACTAACTTG	TGCCTGAACT A	ACCGCAGCCG TGGCGTCGGC	GAGAGCGCCG CTCTCGCGGC	GGCAACTCTG CCGTTGAGAC	
+2	L T V	R V V Q	P N A	T A W	S E A	
451 (GCTCACAGTA CGAGTGTCAT	CGCGTAGTGC A	AACCGAACGC TTGGCTTGCG	GACCGCATGG CTGGCGTACC	TCAGAAGCCG AGTCTTCGGC	

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GTCCGAAAGA AAGTGTCTAC ACCTAACCGC TATTTTTTGT TGACGACTGC +2 P L R D Q F T R A P L D N D I G V	
3201 CCGCTGCGCG ATCAGTTCAC CCGTGCACCG CTGGATAACG ACATTGGCGT GGCGACGCGC TAGTCAAGTG GGCACGTGGC GACCTATTGC TGTAACCGCA	
T+2 S E A T R I D P N A W V E R W K	
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+2 A A G H Y Q A E A A L L Q C T A D	
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+2 T L A D A V L I T T A H A W Q H Q	
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3901 GGGGAAAACC TTATTTATCA GCCGGAAAAC CTACCGGATT GATGGTAGTG CCCCTTTTGG AATAAATAGT CGGCCTTTTG GATGGCCTAA CTACCATCAC	
+2 G Q M A I T V D V E V A S D T P H	
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+2 N W L G L G P Q E N Y P D R L T	
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+2 Y V F P S E N G L R C G T R E L N	
4151 TACGTCTTCC CGAGCGAAAA CGGTCTGCGC TGCGGGACGC GCGAATTGAA ATGCAGAAGG GCTCGCTTTT GCCAGACGCG ACGCCCTGCG CGCTTAACTT	
+2 Y G P H Q W R G D F Q F N I S R	
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TGTCAGTTGT CGTTGACTAC CTTTGGTCGG TAGCGGTAGA CGACGTGCGC +2 E E G T W L N I D G F H M G I G G	
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+2GRYHYQLVWCQKRSDYK	
4401 GTCGCTACCA TTACCAGTTG GTCTGGTGTC AAAAAAGATC TGACTATAAA CAGCGATGGT AATGGTCAAC CAGACCACAG TTTTTTCTAG ACTGATATTT	
+2 D E D L D H H H H H R	
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4551 CACCATATTG CCGTCTTTTG GCAATGTGAG GGCCCGGAAA CCTGGCCCTG GTGGTATAAC GGCAGAAAAC CGTTACACTC CCGGGCCTTT GGACCGGGAC	
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5601	CTGGGTACCG	CTACGGACGA	ACGGCTTATA	GTACCACCTT	AATGGCCGCT TTACCGGCGA
5651		GTAGCTGACA	CCGGCCGACC	CACACCGCCT	CCGCTATCAG GGCGATAGTC
5701	CTGTATCGCA	ACCGATGGGC	ACTATAACGA	CTTCTCGAAC	GCGGCGAATG CGCCGCTTAC

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5751					GATTCGCAGC CTAAGCGTCG	
5801			CTTGACGAGT GAACTGCTCA		GGGACTCTGG CCCTGAGACC	
5851			AAGATTTTAT TTCTAAAATA			
5901			TAGGTTTGGC ATCCAAACCG			
5951			ATACATAACT TATGTATTGA			
6001	GTTCCAGTCC	TTGTCTACCT	ACAGCTGAAT TGTCGACTTA	TACCCGGTTT		
6 0 51			CGGCTCAGGG GCCGAGTCCC			~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
都01 位 上	ACTTATACCC	GGTTTGTCCT	TATCTGTGGT ATAGACACCA	TTCGTCAAGG	ACGGGGCCGA	
6 1 751	CAGGGCCAAG GTCCCGGTTC		CCCCAGATGC GGGGTCTACG			
6 <u>2</u> 01	CTAGAGAACC GATCTCTTGG		TCCAGGGTGC AGGTCCCACG			
6251	CTGTGCCTTA GACACGGAAT					
6501 	GCGCTTCTGC CGCGAAGACG				-	
6351	GGCGCCAGTC CCGCGGTCAG					
6401	AAACCCTCTT TTTGGGAGAA		CCGACTTGTG GGCTGAACAC			
6451	GGTCTCCTCT CCAGAGGAGA		GATGGGCAGT	CGCCCCAGA	AAGTAAGTAC	
6501 	CAGCATGTAT GTCGTACATA	GTTTTAATTA	AACCAAAAAA	AAGAATTCAT	AAATGTAATT	
6551	ATGGCCATAG TACCGGTATC	AACGTAATTA	CTTAGCCGGT	TGCGCGCCCC	TCTCCGCCAA	
6601	TGCGTATTGG ACGCATAACC	GCGAGAAGGC		GTGACTGAGC	GACGCGAGCC	
6651	TCGTTCGGCT AGCAAGCCGA	CGCCGCTCGC	CATAGTCGAG	TGAGTTTCCG		

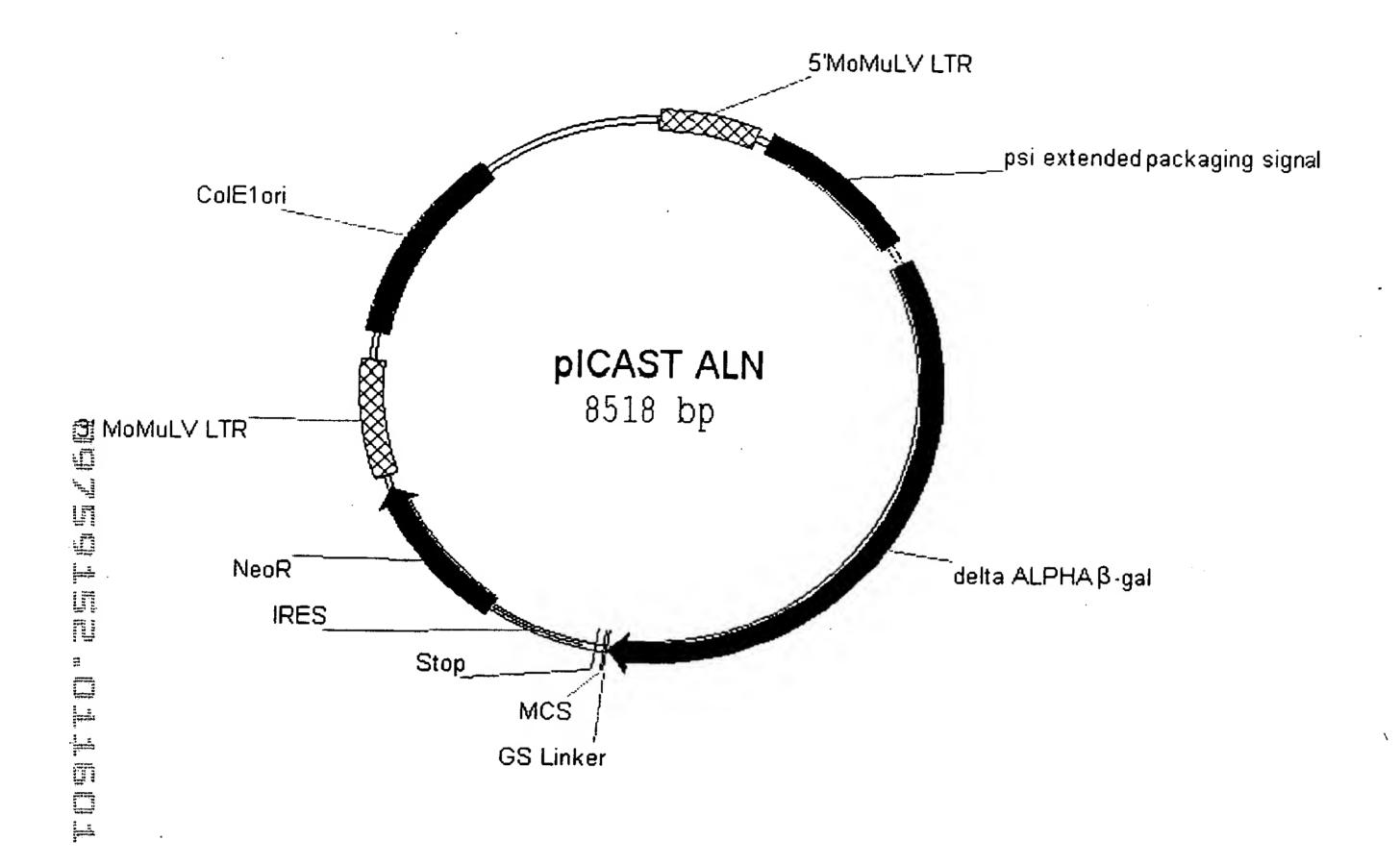


Figure 11A

1	CTGCAGCCTG GACGTCGGAC	AATATGGGCC TTATACCCGG	AAACAGGATA TTTGTCCTAT	TCTGTGGTAA AGACACCATT	GCAGTTCCTG CGTCAAGGAC	
51	CCCCGGCTCA GGGGCCGAGT	GGGCCAAGAA CCCGGTTCTT	CAGATGGAAC GTCTACCTTG	AGCTGAATAT TCGACTTATA	GGGCCAAACA CCCGGTTTGT	
101	GGATATCTGT CCTATAGACA	GGTAAGCAGT CCATTCGTCA	TCCTGCCCCG AGGACGGGGC	GCTCAGGGCC	AAGAACAGAT TTCTTGTCTA	
151	GGTCCCCAGA CCAGGGGTCT	TGCGGTCCAG ACGCCAGGTC	CCCTCAGCAG GGGAGTCGTC	TTTCTAGAGA AAAGATCTCT	ACCATCAGAT TGGTAGTCTA	
201	GTTTCCAGGG CAAAGGTCCC	TGCCCCAAGG ACGGGGTTCC	ACCTGAAATG TGGACTTTAC	ACCCTGTGCC TGGGACACGG	TTATTTGAAC AATAAACTTG	
251	TAACCAATCA ATTGGTTAGT	GTTCGCTTCT CAAGCGAAGA	CGCTTCTGTT GCGAAGACAA	CGCGCGCTTC GCGCGCGAAG	TGCTCCCCGA ACGAGGGGCT	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
301 -11	GCTCAATAAA CGAGTTATTT	AGAGCCCACA TCTCGGGTGT	ACCCCTCACT TGGGGAGTGA	CGGGGCGCCA GCCCCGCGGT	GTCCTCCGAT CAGGAGGCTA	
351 	ACTGACTCAG	CGGGCCCATG	GGCACATAGG	AATAAACCCT TTATTTGGGA	CTTGCAGTTG GAACGTCAAC	
401 TU	CATCCGACTT GTAGGCTGAA	GTGGTCTCGC	TGTTCCTTGG ACAAGGAACC	GAGGGTCTCC CTCCCAGAGG	TCTGAGTGAT	
451	TGACTACCCG ACTGATGGGC	TCAGCGGGGG AGTCGCCCCC	TCTTTCATTT AGAAAGTAAA	GGGGGCTCGT CCCCCGAGCA	CCGGGATCGG GGCCCTAGCC	
\$ 0 1 (T) -(2)	GAGACCCCTG CTCTGGGGAC	CCCAGGGACC GGGTCCCTGG	TGGCTGGGTG	CACCGGGAGG GTGGCCCTCC	GTTCGACCGG	
551 	AGCAACTTAT TCGTTGAATA	CTGTGTCTGT GACACAGACA	CCGATTGTCT GGCTAACAGA	AGTGTCTATG TCACAGATAC	ACTGATTTTA TGACTAAAAT	
601	TGCGCCTGCG ACGCGGACGC	TCGGTACTAG AGCCATGATC	TTAGCTAACT AATCGATTGA	AGCTCTGTAT TCGAGACATA	CTGGCGGACC GACCGCCTGG	
651	CGTGGTGGAA GCACCACCTT	CTGACGAGTT GACTGCTCAA	CTGAACACCC GACTTGTGGG	GGCCGCAACC CCGGCGTTGG	CTGGGAGACG GACCCTCTGC	
701	TCCCAGGGAC AGGGTCCCTG	AAACCCCCGG	CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	
751	CGATGTGGAA GCTACACCTT	TCCGACCCCG AGGCTGGGGC	TCAGGATATG AGTCCTATAC	TGGTTCTGGT ACCAAGACCA	AGGAGACGAG TCCTCTGCTC	• • • • • • • • • • • • • • • • • • • •
801	AACCTAAAAC TTGGATTTTG	AGTTCCCGCC T	TCCGTCTGAA AGGCAGACTT	TTTTTGCTTT AAAAACGAAA	CGGTTTGGAA GCCAAACCTT	
851	CCGAAGCCGC GGCTTCGGCG	GCGTCTTGTC 'CCCAGAACAG	TGCTGCAGCA ACGACGTCGT	TCGTTCTGTG AGCAAGACAC	TTGTCTCTGT AACAGAGACA	
	CTGACTGTGT GACTGACACA	TTCTGTATTT (AAGACATAAA (GTCTGAAAAT CAGACTTTTA	TAGGGCCAGA ATCCCGGTCT	CTGTTACCAC GACAATGGTG	

951 	TCCCTTAAGT AGGGAATTCA		GTAACTGGAA CATTGACCTT			
1001	ACAACCAGTC TGTTGGTCAG		AAGAAGAGAC TTCTTCTCTG			
1051	GCAGAATGGC CGTCTTACCG		CGTCGGATGG GCAGCCTACC			
		TAGTGGGTCC	AATTCTAGTT	CCAGAAAAGT	GGACCGGGCG	
1151	ATGGACACCC TACCTGTGGG		CCCTACATCG GGGATGTAGC			
1201			GTTCGGGAAA			
1251 11			CGTCTCTCCC GCAGAGAGGG	GGAACTTGGA		
1301	CCCCGCCTCG	ATCCTCCCTT	TATCCAGCCC		TCTAGGCGCC	
	GGGGCGGAGC	TAGGAGGGAA	ATAGGTCGGG	AGTGAGGAAG	AGATCCGCGG	
1351	GGCCGCTCTA	GCCCATTAAT	ACGACTCACT	ATAGGGCGAT	TCGAACACCA	
	CCGGCGAGAT	CGGGTAATTA	TGCTGAGTGA	TATCCCGCTA	AGCTTGTGGT	
1 <i>4</i> :01	TGCACCATCA	TCATCATCAC	GTCGACTATA	AAGATGAGGA	CCTCGAGATG	
Separate Sep			CAGCTGATAT			
1451	GGCGTGATTA	CCGATTCACT	GECCETCETE	GCCCGCACCG	ATCCCCCTTC	
			CCGGCAGCAC			
1501	CCAACAGTTA GGTTGTCAAT		ATGGCGAATG TACCGCTTAC			
1551	CACCAGAAGC					
	GTGGTCTTCG	CCACGGCCTT	TCGACCGACC	TCACGCTAGA	AGGACTCCGG	
1601	GATACTGTCG	ずではずでででですで	AAACTGGCAG	атссассетт	ACGATGCGCC	
1001			TTTGACCGTC			
1651	CATCTACACC		ATCCCATTAC TAGGGTAATG			
1701	CCACGGAGAA	TCCGACGGGT	TGTTACTCGC	TCACATTTAA	TGTTGATGAA	
-	GGTGCCTCTT	AGGCTGCCCA	ACAATGAGCG	AGTGTAAATT	ACAACTACTT	
1751	AGCTGGCTAC TCGACCGATG		GACGCGAATT CTGCGCTTAA			· :
	•					
1801	GGCGTTTCAT CCGCAAAGTA				•	•
1851	GTCGTTTGCC CAGCAAACGG					
		CHORCITAM		JIMMMIUC		

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1901	AACCGCCTCG TTGGCGGAGC	GCCACTACCA	GCTGCGCTGG CGACGCGACC	TCACTGCCGT	CAATAGACCT
1951	AGATCAGGAT TCTAGTCCTA				GTCTCGTTGC CAGAGCAACG
2001	TGCATAAACC ACGTATTTGG		ATCAGCGATT TAGTCGCTAA		
2051	AATGATGATT TTACTACTAA				AGATGTGCGG TCTACACGCC
2101	CGAGTTGCGT GCTCAACGCA				CAGGGTGAAA GTCCCACTTT
2151	CGCAGGTCGC GCGTCCAGCG				TATCGATGAG ATAGCTACTC
2 2 01	GCACCACCAA	TACGGCTAGC	GCAGTGTGAT	GCAGACTTGC	TCGAAAACCC AGCTTTTGGG
_	GAAACTGTGG CTTTGACACC				
2 <u>30</u> 1					CGATGTCGGT GCTACAGCCA
2 35 1		ACGCCTAACT	•	GACGACGACT	TGCCGTTCGG
2 10 1 	CAACGACTAA	-			CTGCATGGTC GACGTACCAG
2 4 5 1	AGGTCATGGA				GATGAAGCAG CTACTTCGTC
2501	AACAACTTTA TTGTTGAAAT				ATCCGCTGTG TAGGCGACAC
2551	GTACACGCTG CATGTGCGAC				GAAGCCAATA CTTCGGTTAT
2601		GCCGTACCAC	GGTTACTTAG	CAGACTGGCT	TGATCCGCGC ACTAGGCGCG
		GCTACTCGCT	TGCGCATTGC	GCTTACCACG	AGCGCGATCG TCGCGCTAGC
2701		TCACACTAGT	AGACCAGCGA	CCCCTTACTT	TCAGGCCACG AGTCCGGTGC
2751	GCGCTAATCA CGCGATTAGT	CGACGCGCTG GCTGCGCGAC	TATCGCTGGA ATAGCGACCT	TCAAATCTGT AGTTTAGACA	CGATCCTTCC GCTAGGAAGG
2801				CTGTGGTGCC	CCACCGATAT GGTGGCTATA

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2851				AGACCAGCCC TCTGGTCGGG		
2901	ACGGCTTTAC	CAGGTAGTTT	TTTACCGAAA	CGCTACCTGG GCGATGGACC		
2951				ATGGGTAACA TACCCATTGT		
3001				GTATCCCCGT CATAGGGGCA		
3051				TGATTAAATA ACTAATTTAT	TGATGAAAAC ACTACTTTTG	
3101	CCGTTGGGCA	CCAGCCGAAT	GCCGCCACTA	AAACCGCTAT		
3 1.5 1		ACATACTTGC	CAGACCAGAA	TGCCGACCGC ACGGCTGGCG	TGCGGCGTAG	
32 0 1				AGTTTTTCCA TCAAAAAGGT		
3 25 1	AGGCCCGTTT	GGTAGCTTCA	CTGGTCGCTT	TACCTGTTCC ATGGACAAGG		
3301 				GGATGGTAAG CCTACCATTC		
3 35 1 ————				AAGGTAAACA TTCCATTTGT		·
3 40 1				GGGCAACTCT CCCGTTGAGA		·
3451				GTCAGAAGCC CAGTCTTCGG		
3501				ACCTCAGTGT TGGAGTCACA		·
3551	CGGCGCAGGG	TGCGGTAGGG	CGTAGACTGG	TGGTCGCTTT		
3601 				TAACCGCCAG ATTGGCGGTC		·
3651				AACTGCTGAC TTGACGACTG		
3701	GATCAGTTCA CTAGTCAAGT			GACATTGGCG CTGTAACCGC		
3751	GACCCGCATT CTGGGCGTAA			ACGCTGGAAG TGCGACCTTC		

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3801				GCACGGCAGA CGTGCCGTCT		
3851				TGGCAGCATC ACCGTCGTAG		
3901				TGATGGTAGT ACTACCATCA		
3951				ATACACCGCA TATGTGGCGT		
4001	ATTGGCCTGA TAACCGGACT	ACTGCCAGCT TGACGGTCGA	GGCGCAGGTA CCGCGTCCAT	GCAGAGCGGG CGTCTCGCCC	TAAACTGGCT	
4051				CCGCCTTACT		
4 T 01	AACTGGCGAC	CCTAGACGGT		TGTATACCCC		
4151	CCGAGCGAAA	ACGGTCTGCG TGCCAGACGC	CTGCGGGACG	CGCGAATTGA GCGCTTAACT		·
4201		CGCGGCGACT	TCCAGTTCAA	CATCAGCCGC GTAGTCGGCG		
4 <u>25</u> 1 <u>+</u>	TCGTTGACTA	CCTTTGGTCG		TGCTGCACGC ACGACGTGCG		
4 3 0 1	ACATGCTGA TGTACCGACT			GGGATTGGTG CCCTAACCAC		
4351	CTGGAGCCCG GACCTCGGGC					
4401	ATTACCAGTT TAATGGTCAA			CTGGAGGTGG GACCTCCACC		
4451	CCTTGGCGCG GGAACCGCGC					
4501	ATAAGTGACT TATTCACTGA			GCTGGTTAAG	GCCAATAAAA	
4551	CCACCATATT GGTGGTATAA			CCCGGGCCTT	TGGACCGGGA	
	GTCTTCTTGA CAGAAGAACT	GCTCGTAAGG	ATCCCCAGAA	AGGGGAGAGC	GGTTTCCTTA	
4651	GCAAGGTCTG CGTTCCAGAC	TTGAATGTCG AACTTACAGC	TGAAGGAAGC ACTTCCTTCG	AGTTCCTCTG TCAAGGAGAC	GAAGCTTCTT CTTCGAAGAA	
4701	GAAGACAAAC CTTCTGTTTG	TTGCAGACAT	CGCTGGGAAA	CGTCCGTCGC	CTTGGGGGGT	

4751				•	AAGATACACC TTCTATGTGG
4801				-	ATAGTTGTGG TATCAACACC
4851					GCTGAAGGAT CGACTTCCTA
4901					CTCGGTGCAC GAGCCACGTG
4951	ATGCTTTACA TACGAAATGT				GCCCCCGAA CGGGGGCTT
5001 =		GTGGTTTTCC CACCAAAAGG			
5 05 1					TGGAGAGGCT ACCTCTCCGA
5101 					GATGCCGCCG CTACGGCGGC
5 15 1					CAAGACCGAC GTTCTGGCTG
52 0 1					GGCTATCGTG CCGATAGCAC
5 25 1	CGACCGGTGC	TGCCCGCAAG	GAACGCGTCG	ACACGAGCTG	GTTGTCACTG CAACAGTGAC
5301	TTCGCCCTTC	CCTGACCGAC	GATAACCCGC	TTCACGGCCC	GCAGGATCTC CGTCCTAGAG
5351	GACAGTAGAG	TGGAACGAGG	ACGGCTCTTT	CATAGGTAGT	TGGCTGATGC ACCGACTACG
		GACGTATGCG	AACTAGGCCG	ATGGACGGGT	AAGCTGGTGG
		AGCGTAGCTC	GCTCGTGCAT	GAGCCTACCT	TCGGCCAGAA
		TACTAGACCT	GCTTCTCGTA	GTCCCCGAGC	GCGGTCGGCT
		TCCGAGTTCC	GCGCGTACGG	GCTGCCGCTC	CTAGAGCAGC
	ACTGGGTACC	GCTACGGACG	AACGGCTTAT	AGTACCACCT	TTTACCGGCG
2021	TTTTCTGGAT AAAAGACCTA	AGTAGCTGAC	ACCGGCCGAC	CCACACCGCC	mı .

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5701	GGACATAGCG CCTGTATCGC	TTGGCTACCC AACCGATGGG				
5751	GGGCTGACCG CCCGACTGGC	CTTCCTCGTG GAAGGAGCAC				
5801	CGCATCGCCT GCGTAGCGGA	TCTATCGCCT AGATAGCGGA				
5851	GGGTTCGCAT CCCAAGCGTA	CGATAAAATA GCTATTTTAT				
5901	GGGAATGAAA CCCTTACTTT	GACCCCACCT CTGGGGTGGA				
5951	CATTTTGCAA GTAAAACGTT	GGCATGGAAA CCGTACCTTT				
6001		GAACAGATGG CTTGTCTACC				
6 05 1		AGTTCCTGCC TCAAGGACGG				·
6 10 1		GCCAAACAGG CGGTTTGTCC				
6 1 51		GAACAGATGG CTTGTCTACC				
6201		CATCAGATGT GTAGTCTACA			CTGAAATGAC GACTTTACTG	
6251		ATTTGAACTA TAAACTTGAT			CTTCTGTTCG GAAGACAAGC	
6301	CGCGCTTCTG GCGCGAAGAC				CCCTCACTCG GGGAGTGAGC	·
6351	GGGCGCCAGT CCCGCGGTCA				GTGTATCCAA CACATAGGTT	
6401	ATTTGGGAGA	ACGTCAACGT	AGGCTGAACA	CCAGAGCGAC	TTCCTTGGGA AAGGAACCCT	
6451					TTTCATTCAT AAAGTAAGTA	
6501	GCAGCATGTA CGTCGTACAT				ATTTACATTA TAAATGTAAT	
6551	AATGGCCATA TTACCGGTAT	_		·	GAGAGGCGGT CTCTCCGCCA	
6601	TTGCGTATTG AACGCATAAC				GCTGCGCTCG CGACGCGAGC	* * * - *
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CARTACAGA GARTCAGGG ATAAGCAGGA AAAGAACATG TGACCAAAAG CARTACAGGG CTTAGTCCCC TATTCCCTCC TGCCTTTTC CGGTCGTTT CCGGTCCTTG GCATTTTCC GGCGCACCA CCCCAAAAAG 6801 CARAGGGTCC GCCCCCCTGA GGACCATCAC AAAAATGGAG GCTCAAACAG GTATCCGAGC GCCCCCCTGA GGACCATCAC AAAAATGGAG GCTCAAACAG 6851 GAGGTGGCGA AACCCGACAG GACTATAAAC ATACCAGGCC TTTCCCCTG CTCCACCGCT TTGGGCTGTC CTGATATTC TATGGTCCAC AAAAATGGAC 6901 GAAGCTCCCT CGGGCCCTC CCTGTTCCGA CCCTGCCGCT TACCGGGATCAC CTTCGAGGGA GACCCGACAG GGACAAGGCT GGACTCACAGCC TTTCCCCCTG CTCCACCGCT TTGGGCTGTC CTGATATTC TATGGTCCAC AAAAGGGGGAC 6901 GAAGCTCCCT CGTGGCCTC CCTGTTCCGA CCCTGCCGCT TACCGGGATAC CTTCGAGGGA GACACGGAGA GGACAAGGCT GGACTGCACAC ATAGCTCACG GCACAGCCGGA AAACAGGGAGA CCCCTTCCACC CGGCAAGAGA TAGCCTCACG 7061 CTGTAGGGTT CTCAGTTCGG TGTAGGTCGT TCCCCCAGG CTGGCCTTTCC ACCGGATAC 7061 CTGTAGGGTT CTCAGTTCGG TGTAGGTCGT TCCCCCAGG CTGGCCTTTCC ACCGGATAC 7061 TGCAGGGAAC CCCCGTTCAC CCGGACAGGA TAGCCTCACG GCACACCATA GAGTCAGACC ACACCAGCA AAGGAGGTTC GACCCGGACC 7061 TCGAGGGAAC CCCCGTTCAC CCCGACCCGA CCCGACACACA 7061 TCGAGGGAAC CCCCGTTCAC CCCGACCCGA CCCCACACAAAAGGAGGTTC GGCCTTTTCC CACTCAGAC 7061 TCGAGGGAAC CACCCGGATAC TCCTCCCCAC CGGCACTCATA 7061 CCCTGAGT CCAACCCGGT AAGACAGGAC TTATCGCCAC TGGCAGCACC 60646AACCAA GGGTTAGCA GACCAGGGTA TGTAGGCGCT GCCCTTTAC CGGTCACCT 7061 CCTGGATAC AGGATTAGCA GACCAGGGTA TGTAGGCGCT GCCCTTTCC CGCACACTTCAT CCTAAACCAC 7061 TCTGAACTG TCCCACCGGATAC TCCGCTCCAT AATACCCGCT GCACTCACAC 7061 TCTGAACCT TCCTCAACCC AGTTACCTC GGAAAAAACAACA 7061 TCTGAACCG AAACAAAACCA CCCCTGGTGAC CTAAGGGGT TCTTTTGCT AGACCACTT TCCTCACCCC AGTTACCTCC GGAAAAAACCA 7061 TCTGACCGC AAACAAAACCA CCCCTGGGTG TTTTTCCCCCAC CACTCTCTCT AGACCCCAC AACCAGGAC TTCCTTCTG TCCATAGACAC 7061 TCTGACCGC AAACAAAACCA CCCCTGGGAC CTGCACAAAA AAACAAAACCA 7061 TTGTTACGG GTCTTATGCT TCGCTCCAT ACACCACAAAA AAACAAAACCA 7061 TTTCTTACGG GTCTTATGCT TTTCTTCCTC AGACTCTTT TTCCTTCCAACCACAAAAAAAACAACCT 7061 TTTCTTACGG GTCTTATAGGT TTTTCCTTCCAACCGT TTTTCCTCCAATGCT TTTTCCTCCAATGCT TTTTTTCCTAATGCT TTTTTTCCTAATGCT TTTTTTTCCTAATGCT TTTTTTTCAAACCAA TTTTTCCTTCAATGC	6651		TGCGGCGAGC ACGCCGCTCG	CCATAGTCGA	GTGAGTTTCC		
COGTCATTT COGCTCCTG GCATTTTCC GGGCCAACA CGCAAAAAG 6801 CATAGGCTCC GCCCCCCTCA CGAGCATCAC AAAAATCCAC GCTCAACTCA GTATCCGAGG CGGGGGGACT GCTCGTNGTG TTTTTAGCTG CGAGTTCAGT 6851 GAGGTGGCGA AACCCCACCAGAG GACTATAAAG ATACCAGGC TTTCCCCCTG CTCCACCGCT TTGGGCTGC CCTGATATTC TATGGTCCC AAAGGGGGAC 6901 GAACCTCCT CGTGCGCTCT CCTGTTCCCA CCCTGCGCT TACCGGATAC CTTCGAGGGA GCACCGCAGA GGACAAGGCT GGGACGCGA ATGGCCTATC 6911 CTGTCGCCCT TTCTCCCTTC GGGAAGGCTG GGCTTTCT ATAGCTCAGG GCAGAGGGA AAGGGGGAA GGCCTTCCCAC CCGCAAAGAG TATCCACGG GCACACCATA GAGTCAAGCC ACCCGACA GACCACAGAC TACCCCCAC CTGCGCTCTC GCGCAAGACC CCCCGTTCAG CCCGACCGCACA GACCACACAC 100 CTGTAGGTAT CTCACTTCCG TCTACCCCAC CCGCAAGAGA TATCCACTCC GCGCAACACC CCCCGTTCAG CCCGACCGCA CGCGAAGAGA TATCCACTCAC GCACACCCCTTCAGC CCCGCTCCAC CGCGACACACAC 101 CCCCTTGAGCT CCAACCCGC AAGACCACACCAC CGCGCACACA 102 CCCCTTCAGCT CCAACCCGC AAGACCACAC CTACCCCAC CGCCCACAC 103 CCCCCACACC CCCCGTTCAG CCCGACCGCT GCGCCTATC CGCTCACAC 104 CCCCTTGAGCT CCAACCCGC AAGACCACAC CTACCCCCAC CGCACACAC 105 CCCCAACACC CCCCGTTCAG CCCGACCACAC 105 CCCCAACACC CCCCGTTCAG CCCGACCACAC 106 CCCCAACCAC CCCGCTTCAG CCCGACCCACAC 107 CCCCAACCCAC CCGCTCACACCCAC CACCACACACAC 107 CCCCAACCCCAC CCGCTCACACCCAC CACCACACACAC 107 CCCCAACCCAC CCGCGATCA CTCACCCCAC CCACCACACAC 107 CTTCACACCCCC CCCCTCCAC CTCACACCCAC CCACCACACAC 107 CTTCACACCCCC CCCCCTCCAC CTCACACCACAC CACCACCACAC 107 CTTCACACCCCC CCCCCTCCAC CCCACCACAAAAAACACCT 107 CTTCACCCCC CACCCGAACCC ACTTACCCTC CCCACCACAAAAAAACACCT 107 CTCCCACCCCCACCCGACCCCACCACCACACACACCACCACC	6701						
6801 CATAGGETC GCCCCCTGA CGACCATCAC AAAAATCGAC GCTCAAGTCA GTATCCGAGG CGGGGGGGACT GCTCCACCGA AACCCGACAG GACTATAAAG ATACCAGGGC TTTCCCCCTG CTCCACCGCT TTGGGCTGC CGACTATATAC ATAGCCAGGA CCCTCCACCGCT TTGGGCTGC CGACTATACAC CCTCCACCGCT TTGGGCTGC CGACTATACAC CCTCCACCGCT TTGGGCTGC CGACAGGAGGAC CCTCCACCGCT TACCGGATAC CCTCCACCGC TTCCCACCGA GACCAGGAG GACCAGGAG GGACCAGGCT GGGCCTGCAC ATAGCCTATG 6751 CTGCAGGCA CACCCGAG GGACCAGGGG GGGCTTCCC ATAGCTCACG 40 GACCACCCAT CTCCCCTC GGGAAGGGT GGGCTTTCC ATAGCTCACG 41 CCTCTAGGTAC CCACCCTCGCAC CCCGCAGAGAGA TACCGCTATG 42 CCTCTAGGTAC CTCAGTTCGG TCTAGCTCCT TCCCTCCAG CTGGCCTCTG 43 CACCCCTTCGAC CCCCCTTCCAC CCCGCAAGAGA TACCGACTGC 44 CCTCTAGGTAC CACCCGGT AGCCACCGT GCCCCTTACC GACCCGACA 45 CCTCTTGGCTAC CAACCCGGT AGCCACCGT GCCCCTTAC CGGCACCAC 46 CCTCTTGGCTAC CAACCCGGT AGCCACGCT TTTCCCCCAC TGCCACCACC 47 CCTCTAGGTAC CAACCCGGT AGCCACGCT TATCGCCCAC TGCACCACC 47 CCTCTAGACT CCAACCCGGT AGCCACGCT TATCGCCCAC TGCACCACC 47 CCTCTTCAACT CCAACCCGGT AAACCACGAC TTATCGCCCAC TGCACCACGC 47 CCTCTTCAACT CCAACCCGGT AAACCACGAC TATCGCCCAC CGAATATGCTCC 47 CCTCTTCAACT CCAACCCGGT AACCACGAC TATCGCCCAC CGATCTCCAC 47 CTCTCAACCATTC TCCTAATCGT CTCGCTCCAT ACATCCCCCC CGATCTCCA 47 CTCTCAACCATTC TCCTAATCGT CTCGCTCCAT ACATCCCCCC CGATCTCCA 47 CTCTCAACCATTC TCCTAATCGT CTCGCTCCAT ACATCCCCCC CGATCTCCA 47 CTCTCAACCATC TCCTCAACCCC ACTTTCTCT CAACCACCGC 47 CTCTCAACCATC TCCTCAACCCC TCCATCCACCAC CACCACACAAC ACTATTTGT 47 CTCTCAACCATC TCCTCAACCC ACTTACCTCC CGAAAAAACCA 47 CTCTCACCCCC TTCTTTTTCT TCCATCAACCAC CCTCTTTTCTC TCCATAAACCT 47 CTCTCCCCCC TTCTTTTTCTT TCCTTCAACCAC CCTCTTTTCTC TCCATAAACCT 47 CTCTCCCCCC TCCTCAACCCC TCCATCAAAAACCA CCTCTTTTTCCT AACCATCCAC 47 CTCTCCCCCC TCCTCAACCCC TCCCTCCAACAAAAAACCA CCTCTTTTTCCTAACCATC CTTTTTCCTAACCATC CAACCACCACAAAAAACCA CCCCTCTCCATCCA	6751		CCGGTCCTTG	GCATTTTTCC	GGCGCAACGA	CCGCAAAAAG	
CTCCACCGCT TIGGGCTGT CTGATATITC TATGGTCCGA AAAGGGGAC 6901 GAAGCTCCT CGTGCGCTCT CCTGTCCGA CCCTGCCGCT TACCGGATAC CTTCAGAGGGA GCACGGGACA GGACAGGCT GGGACGGCCA ATGGCCTATG 6751 CTGTCCGCCT TICTCCCTTC GGGAAGGCT GGGACGGCCA ATGGCCTATG GACACCGCCA AAGAGGGAAG CCCTTGCGCC CGCGAAAGGA TATGCGTCACG GACACCCATA GAGTCAAGC CCCCGTTCGAC CGCGAAAGGA CTGCGCGCA 10 CTGTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG GACACCCATA GAGTCAAGC ACATCCAGCA AGCCAGCATA GCCCGACAGC 10 CTGTAGGTAC CCCAGTCAG CCCGACCGCT GCGCCTTATC CGGTAACTAT ACGTGCTTGG GGGCCAAGTC GGCCTGCGCA CCCGGAATAG GCCATTGATA 11 CACTGGTAC GGTTGGGCCA TTCTGTGCGCA TGGCACAGCAGC GCAGAACTCA GGTTGGGCCA TTCTGTGCGCA CTGGACACAGC GCAGAACTCA GGTTGGGCCA TTCTGTGCCG ACACAGCAGC GCAGAACTCA GGTTGGGCCA TTCTGTGCCCA CAATAGCGCCC CGATGGTCCA 12 CACTGGTAC AGGATTAGCA GACCAGCGAT TACAGCACCA CGATGATACAGCACCACAGCACAG	6801		GCCCCCTGA	CGAGCATCAC	AAAAATCGAC	GCTCAAGTCA	
CTTCGAGGGA GCACCGAGA GGACAGGCT GGGACGGCA ATGGCTATG 6351 CTGTCCGCCT TTCTCCCTC GGGAGGCGG GCGCTTTCTC ATAGCTCACG GACAGGCGGA AAGAGGGAGA CCCTTCGCAC CGCGAAAGAG TATCGAGTGC 7051 CTGTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG GACATCCATA GAGTCAAGCC ACATCCAGCA AGCGAGGTTC GACCCGACAC ACCTGCTTGAG GGGGCAAGCC GCCGACCGGT GGCCTTATC CGGCTACTAT ACCTGCTTGG GGGGCAAGCC GCCGACCGGT CGCGCAATAG GCCATTGATA 7191 CGTCTTGAGT CCAACCCGGT AAGACAGGAC TTATCGCCAC TGGCAGCAC CGAGAACTCA GGTTGGGCCA TTCTGTGCTG AATAGCGGT ACCTCCAGC GCAGAACTCA GGTTGGGCCA TTCTGTGCTG AATAGCGGT ACCTCCAGC GCAGAACTCA GGTTGGCCCA TCCTGCTCCAT ACACCCGCC AGATGCTCA 7101 CTTGAAGTG GTGGCCTAAC TACGGCTACA CTACAAGAACA AGTATTTGGT AGAACTTCAC CACCGGATTG ATGCCGATGT GATCTTCTTT TCATAAACCA 7251 ATCTGCGCTC TGCTGAACC AGTTACCTTC GGAAAAAAGGA TTGTATAGCCA 7301 TTGATCCGGC AAACAAACCA CCGCTGGTAG CGGTGCTTT TTTGTTTCCA AACACTGCAG ACGACTTTGGT GGCGACCACC GCCACAAAA AAACAAACGT 7351 AGCAGCAGAT TACGGGCAGA AAAAAAAGGA TCCTATTATCC AACCATCGAG 7401 TTTTCTCACG GGTCTGACGC CAGCGGGGCAC GAGACCACAAAA AAACAAACCT 7401 TTTTCTCACG GGTCTGACGC CAGCAGGCAC GAGACCTCTTTCTC AACCATCGAG 7401 TTTTCTCACG GGTCTGACGC CAGCAGCAC GAGCACCTCTTTCT AGCAACACAG 7501 CGCAAATCAA ACCGGCTCT TTTTTCCTA GAGTCCTTC AGCAACCAG 7501 CGCAAATCAA TCCTAAAGTAT ATATCAGTAA ACTTGGTCCT ACGATTACCC 7501 CGCAAATCAA TCTAAAGTAT ATATCAGTAA ACTTGGTCTC ACCATTACCA GCGTTTAGT AGATTTCATA TATATCACTAT TGAACCCAGC TGTCTCCAT 7551 ATGCTTAATC AGTGAGGCAC CTACTCCACT TTTCGGTCA ACCTTACCCTA 7551 ATGCTTAATC AGTGAGGCAC CTACTCCACT TTTCGGTCA ACCTTACCCA GCGTTTAGT AGATTTCATA TATATCACTAT TGAACCCAGC TGTCTCCAT 7551 ATGCTTAATC AGTGAGGCAC CTACTCCACC CATCCTCCACAC TTTTCCCTCACCACTCCACACC 7551 ATGCTTAATCA AGTGAGCAC CTACTCCACC CTTTTCCCTC 7551 ATGCTTAATC AGTGAGGCAC CTACTCCACC CTTCTCCACC CTTTTCCCTC 7551 ATGCTTAATC AGTGAGGCAC CTACTCCACC CTTCTCCACC CTTCTCCACC 7551 ATGCTTAATC AGTGAGCAC CTACTCCACC CTTCCTCCAC CTTCTCCACC 7551 ATGCTTAATC AGTGAGCAC CTACTCCACC CTTCCTCCAC CTTCTCCACC 7551 ATGCTTAATC AGTGAGCAC CTACTCCACC CTTCCTCCACC CTTCCCACCACACACA	6851						
GACAGGGGA AAGAGGGAAG CCCTTCGCAC CGCGAAAGAG TATCGAGTGC TOTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG GACATCCATA GAGTCAAGCC ACATCCAGCA AGCAGGTTC GACCCGACAC TOTAGAGTAT CTCAGTTCG GGGCACGCT GGCGCTTATC CGGTAACTAT CGCAGAACC CCCCGTTCAG CCCGACCGCT GGCGCGTATC CGGTAACTAT CACTGCTTCG GGGGCAAGTC GGGCTGGCGA CGCGGAATAG GCCATTGATA TOTAGACGAC CCAACCCGGT AAGACACGAC TTATCGCCAC TGGCAGCACC GCAGAACTCA GGTTGGGCCA TACGACACGAC TTATCGCCAC TGGCAGCACC GCAGAACTCA GGTTGGGCCA TACGACCGCA CATCGCCAC CGATGTCTCA TOTAGACCACTA GGTTGAAC AGGATTAGCA GAGCAGGAT TGTAGGCGGT GCTACAGAGT GTGACCCATTG TCCTAATCGT CTCGCTCCAT ACATCCGCCA CGATGTCTCA TOTAGAACTCAC CACCGGATTG ATGCCGATGT GATCTCTTG TCATAAACCA TOTAGACCACTA CTCGAACCCAC CACCGGATG ACACCACCACACACACACACACACACACACACACACAC	6901				·		
GACATCCATA GAGTCAAGCC ACATCCAGCA AGGAGGTTC GACCCGACCC							
ACGTGCTTGG GGGGCAAGTC GGGCTGGCGA CGCGGAATAG GCCATTGATA 1	*****					• - •	
7:51 CGCCACACC CAACCCGGT AAGACACGAC TTATCGCCAC TGGCACCACC GCAGAACTCA GGTTGGCCA TTCTGTGCTG AATAGCGGT ACCGTCGTCG 7:51 CACTGGTAAC AGGATTAGCA GAGCGAGGTA TGTAGGCGGT GCTACAGAGT GTGACCATTG TCCTAATCGT CTCGCTCCAT ACATCCGCCA CGATGTCTCA 7:201 TCTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGAAC AGTATTTGGT AGAACTTCAC CACCGGATTG ATGCCGATGT GATCTTCTT TCATAAACCA 7:251 ATCTGCGGCTC TGCTGAAGCC AGTTACCTTC GGAAAAAGAG TTGGTAGCTC TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTTCTC AACCATCGAG 7:301 TTGATCCGGC AAACAAACCA CCGCTGGTAG CGGTGGTTT TTTGTTTGCA AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACCT 7:351 AGCAGCAGAT TACGCGCAGA AAAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCCTCGTCTA ATGCGCGCTC TTTTTTCCTA GAGTTCTCTT AGGAAACTAG 7:401 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGATC CACTTAGGTG CAATTCCCTA 7:451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACCCG 7:501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTTCATA TATACCTATT TGAACCAGAC TGTCAATGGT 7:551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT							
TOTTGAAGTG GTGGCCTAAC TACGGCTCCAT ACATCCGCCA CGATGTCTCA TOTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGAAC AGTATTTGGT AGAACTTCAC CACCGGATTG ATGCCGATGT GATCTTCTTG TCATAAACCA TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTCTC AACCATCGAG TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTTCTC AACCATCGAG TTGATCCGGC AAACAAACCA CCGCTGGTAG CGGTGGTTTT TTTGTTTGCA AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACGT AGCAGCAGAT TACGCGCAGA AAAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCGTCGTCTA ATGCGGCAGA AAAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCGTCGTCTA ATGCGGCACA AACAAACCA CGATGGAAC CAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA TTTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACCCG TOTTGGTCATA AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAA CTTTGCTCAA GCGTTTAGTT AGATTTCATA TATACTCATT TGAACCAGAC TGTCAATGGT ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	71 <u>0</u> 1						
7201 TCTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGAAC AGTATTTGGT AGAACTTCAC CACCGGATTG ATGCCGATGT GATCTTCTTG TCATAAACCA 7251 ATCTGCGCTC TGCTGAAGCC AGTTACCTTC GGAAAAAGAG TTGGTAGCTC TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTTCTC AACCATCGAG 7301 TTGATCCGGC AAACAAACCA CCGCTGGTAG CGGTGGTTTT TTTGTTTGCA AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACGT 7351 AGCAGCAGAT TACGCGCAGA AAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCGTCGTCTA ATGCGCGTCT TTTTTTCCTA GAGTTCTCTT AGGAAACTAG 7401 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA 7451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACCCG 7501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7 15 1		·				
TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTTCTC AACCATCGAG 7301 TTGATCCGGC AAACAACCA CCGCTGGTAG CGGTGGTTTT TTTGTTTGCA AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACGT 7351 AGCAGCAGAT TACGCGCAGA AAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCGTCGTCTA ATGCGCGTCT TTTTTCCTA GAGTTCTCT AGGAAACTAG 7401 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA 7451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACGCCG 7501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7201 						
AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACGT 7351 AGCAGCAGAT TACGCGCAGA AAAAAAGGAT CTCAAGAAGA TCCTTTGATC TCGTCGTCTA ATGCGCGTCT TTTTTTCCTA GAGTTCTTCT AGGAAACTAG 7401 TTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA 7451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACGCCG 7501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTCCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7251						
TCGTCGTCTA ATGCGCGTCT TTTTTCCTA GAGTTCTTCT AGGAAACTAG 7401 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA 7451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACGCCG 7501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7301				_		
AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTTGAGTG CAATTCCCTA 7451 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTGCGGC AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACGCCG 7501 CGCAAATCAA TCTAAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7351				_		
AAACCAGTAC TCTAATAGTT TTTCCTAGAA GTGGATCTAG GAAAACGCCG 7501 CGCAAATCAA TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA GCGTTTAGTT AGATTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7401						
GCGTTTAGTT AGATTTCATA TATACTCATT TGAACCAGAC TGTCAATGGT 7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7451 						
7551 ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT	7501 	GCGTTTAGTT	AGATTTCATA	TATACTCATT	TGAACCAGAC	TGTCAATGGT	
~	7551	ATGCTTAATC	AGTGAGGCAC	CTATCTCAGC	GATCTGTCTA	TTTCGTTCAT	

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7601		CTGACTCCCC GACTGAGGGG				
7651		GCCCCAGTGC CGGGGTCACG				
7701		TTATCAGCAA AATAGTCGTT				
7751		TGCAACTTTA ACGTTGAAAT				
7801	GCCCTTCGAT	GAGTAAGTAG CTCATTCATC	AAGCGGTCAA	TTATCAAACG		
7851		ACAGGCATCG TGTCCGTAGC				
79 01 监		CGGTTCCCAA GCCAAGGGTT				
7951 ① ———	TTGTGCAAAA AACACGTTTT	AAGCGGTTAG TTCGCCAATC				
8901 Tu		GCAGTGTTAT CGTCACAATA				
8051	GAGAATGACA	CATGCCATCC GTACGGTAGG	CATTCTACGA	AAAGACACTG	ACCACTCATG	
8101 111	AGTTGGTTCA	CATTCTGAGA GTAAGACTCT	TATCACATAC	GCCGCTGGCT	CAACGAGAAC	
8 <u>1</u> 51		TATGCCCTAT	TATGGCGCGG	TGTATCGTCT	TGAAATTTTC	
8201 8251		ACCTTTTGCA	AGAAGCCCCG	CTTTTGAGAG	TTCCTAGAAT	
	CCGCTGTTGA GGCGACAACT TTCAGCATCT	CTAGGTCAAG	CTACATTGGG	TGAGCACGTG	GGTTGACTAG	
	AAGTCGTAGA	AAATGAAAGT	GGTCGCAAAG	ACCCACTCGT	TTTTGTCCTT	
8401	CCGTTTTACG	GCGTTTTTTC	CCTTATTCCC	GCTGTGCCTT	TACAACTTAT	
8451	GAGTATGAGA	AGGAAAAAGT.	TATAATAACT	TCGTAAATAG	TCCCAATAAC	
8501	AGAGTACTCG	CCTATGTATA	AACTTACATA	AATCTTTTTA	TTTGTTTATC	
	CCCAAGGCGC					

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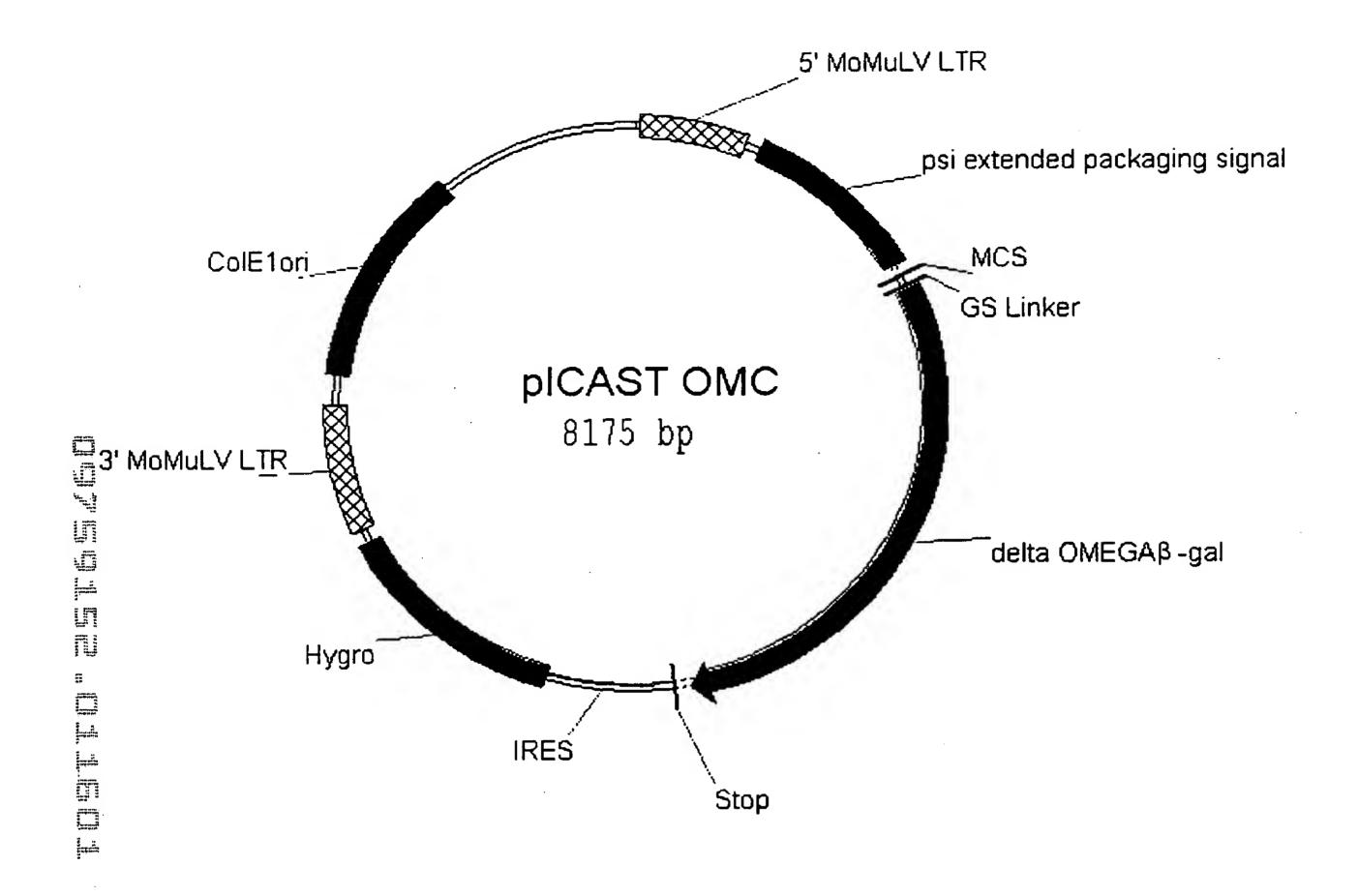


Figure 12A

1	CTGCAGCCTG GACGTCGGAC				-	
51	CCCCGGCTCA GGGGCCGAGT					
101	GGATATCTGT CCTATAGACA					
151	GGTCCCCAGA CCAGGGGTCT	TGCGGTCCAG ACGCCAGGTC				
201	GTTTCCAGGG CAAAGGTCCC	TGCCCCAAGG ACGGGGTTCC				
251	TAACCAATCA ATTGGTTAGT	GTTCGCTTCT CAAGCGAAGA				
301 	GCTCAATAAA CGAGTTATTT					
351 <u>H</u>	TGACTGAGTC ACTGACTCAG				CTTGCAGTTG GAACGTCAAC	
#01 TU	CATCCGACTT GTAGGCTGAA			•		
43 1	TGACTACCCG ACTGATGGGC					· · · · · · · · · · · · · · · · · · ·
501 	GAGACCCCTG CTCTGGGGAC			•		
551	AGCAACTTAT TCGTTGAATA	CTGTGTCTGT GACACAGACA				
601	TGCGCCTGCG ACGCGGACGC				CTGGCGGACC GACCGCCTGG	
651					CTGGGAGACG GACCCTCTGC	
701			CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	
751		AGGCTGGGGC	AGTCCTATAC	ACCAAGACCA	AGGAGACGAG TCCTCTGCTC	
801	AACCTAAAAC TTGGATTTTG				CGGTTTGGAA GCCAAACCTT	
851		CGCAGAACAG	ACGACGTCGT	AGCAAGACAC	AACAGAGACA	
901	CTGACTGTGT GACTGACACA				CTGTTACCAC GACAATGGTG	

951		TTGACCTTAG AACTGGAATC			
1001		GGTAGATGTC CCATCTACAG			
1051	GCAGAATGGC CGTCTTACCG	CAACCTTTAA GTTGGAAATT			
1101	CCGAGACCTC GGCTCTGGAG	ATCACCCAGG TAGTGGGTCC			
1151	ATGGACACCC TACCTGTGGG	AGACCAGGTC TCTGGTCCAG			
1201		CTCCCTGGGT GAGGGACCCA			
1251		CCATCCGCCC			
1301	GGGGCGGAGC	ATCCTCCCTT TAGGAGGGAA	ATAGGTCGGG	AGTGAGGAAG	
1351	GGCCGCTCTA	GCCCATTAAT CGGGTAATTA	ACGACTCACT	ATAGGGCGAT	TCGAATCAGG
1401	GGAACCGCGC	CCGGATCCTT GGCCTAGGAA	TTAATTCGCG	TTAACCCTCC	· · · · · · · · · · · · · · · · · · ·
1451		GCGTGATTAC CGCACTAATG	•		
1501		AACCCTGGCG TTGGGACCGC			
1551	CCCCTTTCGC GGGGAAAGCG	CAGCTGGCGT GTCGACCGCA			
1601	TCCCAACAGT AGGGTTGTCA	TACGCAGCCT ATGCGTCGGA			
1651	GGCACCAGAA CCGTGGTCTT	GCGGTGCCGG CGCCACGGCC			
1701	CCGATACTGT GGCTATGACA	CGTCGTCCCC GCAGCAGGGG			
1751	CCCATCTACA GGGTAGATGT	CCAACGTGAC GGTTGCACTG			
1801		TTAGGCTGCC	CAACAATGAG	CGAGTGTAAA	AATGTTGATG TTACAACTAC
1851	AAAGCTGGCT TTTCGACCGA	TGTCCTTCCG	GTCTGCGCTT	AATAAAAACT	

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1901	TCGGCGTTTC		CANCGGGCGC			
1951	CAGTCGTTTG GTCAGCAAAC		TTGACCTGAG AACTGGACTC			
2001	AAAACCGCCT TTTTGGCGGA		GTGCTGCGCT CACGACGCGA			
2051	GAAGATCAGG CTTCTAGTCC		GATGAGCGGC CTACTCGCCG			
2101			AAATCAGCGA TTTAGTCGCT			
2151	TTAATGATGA AATTACTACT		GCTGTACTGG CGACATGACC			
2201			ACGGGTAACA TGCCCATTGT		CCGTCCCACT	
2251			CCGCGCCTTT	CGGCGGTGAA	ATTATCGATG	·
- - 23 01 TJ			CGCGTCACAC GCGCAGTGTG			
2351			AATCCCGAAT TTAGGGCTTA			
24 01			CGCTGATTGA GCGACTAACT			
2451			GAAAATGGTC CTTTTACCAG			
2501			TAACCGTCAC ATTGGCAGTG			
2551			CGATGGTGCA GCTACCACGT			·
2601	AGAACAACTT TCTTGTTGAA		CGCTGTTCGC GCGACAAGCG			
2651			CTACGGCCTG GATGCCGGAC			
2701	ATAACTTTGG	GTGCCGTACC	TGCCAATGAA ACGGTTACTT	AGCAGACTGG		
2751	GCTGGCTACC CGACCGATGG	GGCGATGAGC CCGCTACTCG	GAACGCGTAA CTTGCGCATT	CGCGAATGGT GCGCTTACCA	GCAGCGCGAT CGTCGCGCTA	* * * * * * * * * * * * * * * * * * *
2801	CGTAATCACC	CGAGTGTGAT	CATCTGGTCG GTAGACCAGC	CTGGGGAATG	AATCAGGCCA	

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2851		CACGACGCGC GTGCTGCGCG				
2901	CCCGCCCGGT GGGCGGGCCA	GCAGTATGAA CGTCATACTT				
2951		CGATGTACGC GCTACATGCG				
3001		TGGTCCATCA ACCAGGTAGT				
3051		CCTTTGCGAA GGAAACGCTT				
3101		AATACTGGCA TTATGACCGT				
3 15 1		TGGGACTGGG ACCCTGACCC				
3 20 1	TGCCGTTGGG	GTGGTCGGCT CACCAGCCGA	ATGCCGCCAC	TAAAACCGCT	ATGCGGCTTG	
32 <u>5</u> 1		TCTGTATGAA AGACATACTT			· · · • • • · ·	
3301		ACGGAAGCAA TGCCTTCGTT				
3351 		AACCATCGAA TTGGTAGCTT				
34 0 1		TCCTGCACTG AGGACGTGAC				
3451	AAGCGGTGAA TTCGCCACTT	GTGCCTCTGG CACGGAGACC			-	
3501		ACTACCGCAG TGATGGCGTC				
3551	GTACGCGTAG CATGCGCATC	TGCAACCGAA ACGTTGGCTT				
3601	GTCGCGGACC	GTCGTCACCG	CAGACCGCCT	TTTGGAGTCA	CACTGCGAGG	
3651		CCACGCCATC GGTGCGGTAG				·
3701		ACCCATTATT	CGCAACCGTT	AAATTGGCGG		
3751		TACACCTAAC	CGCTATTTTT	TGTTGACGAC		

3801		GTGGGCACAG	CTATCTAGAC	AACAGAAACT TTGTCTTTGA	GTAAAGGCTT	
3851				CACCGGTAAT GTGGCCATTA		
3901				ATCCCTCGAC TAGGGAGCTG		
3951	TATTTTCCAC ATAAAAGGTG			ATGTGAGGGC TACACTCCCG		
4001				GGTCTTTCCC CCAGAAAGGG		
	AGGAATGCAA TCCTTACGTT			GGAAGCAGTT CCTTCGTCAA		
4101		TGTTTGTTGC	AGACATCGCT	CCCTTTGCAG GGGAAACGTC		
4 1 51		GCGACAGGTG CGCTGTCCAC	CCTCTGCGGC GGAGACGCCG		CACATATTCT	
4 <u>2</u> 01	TACACCTGCA ATGTGGACGT	AAGGCGGCAC	AACCCCAGTG TTGGGGTCAC	CCACGTTGTG GGTGCAACAC	AGTTGGATAG	
4251	TTGTGGAAAG AACACCTTTC	AGTCAAATGG	CTCTCCTCAA			
4301 1	AAGGATGCCC TTCCTACGGG			GGATCTGATC CCTAGACTAG		
4351	GTGCACATGC CACGTGTACG					
4401	CCCGAACCAC GGGCTTGGTG					
4451	ATGAAAAAGC TACTTTTTCG					
		TCGCAGAGGC	TGGACTACGT	CGAGAGCCTC	CCGCTTCTTA	*
	CTCGTGCTTT GAGCACGAAA	CAGCTTCGAT GTCGAAGCTA	GTAGGAGGGC CATCCTCCCG	GTGGATATGT CACCTATACA	CCTGCGGGTA GGACGCCCAT	
	AATAGCTGCG TTATCGACGC	GGCTACCAAA	GATGTTTCTA	GCAATACAAA	TAGCCGTGAA	
	TGCATCGGCC ACGTAGCCGG	GCGCTCCCGA CGCGAGGGCT	TTCCGGAAGT AAGGCCTTCA	GCTTGACATT CGAACTGTAA	GGGGAATTTA CCCCTTAAAT	
4701	GCGAGAGCCT CGCTCTCGGA	GACCTATTGC CTGGATAACG	ATCTCCCGCC TAGAGGGCGG	GTGCACAGGG CACGTGTCCC	TGTCACGTTG ACAGTGCAAC	

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4751	CAAGACCTGC GTTCTGGACG	CTGAAACCGA GACTTTGGCT				
4801		GCGATCGCTG CGCTAGCGAC				
4851		ACCGCAAGGA TGGCGTTCCT				·
4901	ATATGCGCGA TATACGCGCT	TTGCTGATCC AACGACTAGG	GGTACACATA			
4951 .	CGACACCGTC GCTGTGGCAG	AGTGCGTCCG TCACGCAGGC				·
5001		CTGCCCGAA GACGGGGCTT				
5051		TCCTGACGGA AGGACTGCCT				
- <u> </u> -		ATGTTCGGGG TACAAGCCCC				
-UT 5151 :					CTACTTCGAG GATGAAGCTC .	
5201		CGGAGCTTGC GCCTCGAACG			CGTATATGCT GCATATACGA	
5251	GGCGTAACCA		AGATAGTCTC		GGCAATTTCG CCGTTAAAGC	
5301	ATGATGCAGC		GGTCGATGCG		CCGATCCGGA GGCTAGGCCT	
5351					CGGCCGTCTG GCCGGCAGAC	
5401					CGACGCCCCA GCTGCGGGGT	
5451	CGTGAGCAGG	CTCCCGTTTC	CTTATCTCAT	CTACGGCTGG	GGGATCTATC CCCTAGATAG	
5501	CTATTTTATT	TTCTAAAATA	AATCAGAGGT	CTTTTTCCCC	GGAATGAAAG CCTTACTTTC	
5551	ACCCCACCTG TGGGGTGGAC	ATCCAAACCG	TTCGATCGAA	TTCATTGCGG	TAAAACGTTC	
5601	CGTACCTTTT	ATACATAACT TATGTATTGA	GAGAATAGAG CTCTTATCTC	AAGTTCAGAT TTCAAGTCTA	CAAGGTCAGG GTTCCAGTCC	
5651	AACAGATGGA	ACAGCTGAAT	ATGGGCCAAA	CAGGATATCT	GTGGTAAGCA CACCATTCGT	

5701				ATGGAACAGC TACCTTGTCG		
5751		ATAGACACCA	TTCGTCAAGG	ACGGGGCCGA	GTCCCGGTTC	
5801	AACAGATGGT	CCCCAGATGC	GGTCCAGCCC		CTAGAGAACC	
5851	ATCAGATGTT TAGTCTACAA			TGAAATGACC ACTTTACTGG		.
5901				TTCTGTTCGC AAGACAAGCG		
5951				CCTCACTCGG GGAGTGAGCC		
6001 LT -		GACTCAGCGG	GCCCATGGGC	TGTATCCAAT ACATAGGTTA	TTTGGGAGAA	
6 © 51	CGTCAACGTA		CAGAGCGACA	TCCTTGGGAG AGGAACCCTC		
년 - 6 四 1	GAGTGATTGA CTCACTAACT	CTACCCGTCA GATGGGCAGT	GCGGGGGTCT CGCCCCAGA	TTCATTCATG AAGTAAGTAC	GTCGTACATA	
6 <u>15</u> 1	CAAAATTAAT	TTGGTTTTTT	TTCTTAAGTA	TTTACATTAA AAATGTAATT	ATGGCCATAG	·
6201				AGAGGCGGTT TCTCCGCCAA		
6251				CTGCGCTCGG GACGCGAGCC		
6301				GGTAATACGG CCATTATGCC		
6351		•		GAGCAAAAGG CTCGTTTTCC		·
6401	GCCAGGAACC CGGTCCTTGG		GCGCAACGAC	CGCAAAAAGG	TATCCGAGGC	
		CTCGTAGTGT	AAAATCGACG TTTTAGCTGC	CTCAAGTCAG GAGTTCAGTC	AGGTGGCGAA TCCACCGCTT	
	ACCCGACAGG TGGGCTGTCC	ACTATAAAGA TGATATTTCT	TACCAGGCGT ATGGTCCGCA	TTCCCCCTGG AAGGGGGACC	AAGCTCCCTC TTCGAGGGAG	
6551	GTGCGCTCTC CACGCGAGAG	CTGTTCCGAC GACAAGGCTG	CCTGCCGCTT GGACGGCGAA	ACCGGATACC TGGCCTATGG	TGTCCGCCTT ACAGGCGGAA	
6601		GGAAGCGTGG	CGCTTTCTCA	TAGCTCACGC	TGTAGGTATC	

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6651	AGTCAAGCCA	GTAGGTCGTT CATCCAGCAA	GCGAGGTTCG	ACCCGACACA	CGTGCTTGGG	
6701	CCCGTTCAGC	CCGACCGCTG GGCTGGCGAC	CGCCTTATCC	GGTAACTATC	GTCTTGAGTC	
6751		AGACACGACT TCTGTGCTGA				
6801		AGCGAGGTAT TCGCTCCATA				
6851		ACGGCTACAC TGCCGATGTG				
6901 		GTTACCTTCG CAATGGAAGC	CTTTTTCTCA	ACCATCGAGA		
695 i	AACAAACCAC TTGTTTGGTG	CGCTGGTAGC GCGACCATCG				
	ACGCGCAGAA TGCGCGTCTT	TTTTTCCTAG	AGTTCTTCTA	GGAAACTAGA	TTTCTACGGG AAAGATGCCC	
	GTCTGACGCT CAGACTGCGA	CAGTGGAACG GTCACCTTGC	AAAACTCACG TTTTGAGTGC	TTAAGGGATT AATTCCCTAA	TTGGTCATGA	
710	GATTATCAAA		ACCTAGATCC	TTTTAAATTA	AAAATGAAGT	
715	TTGCGGCCGC AACGCCGGCG	AAATCAATCT TTTAGTTAGA				
7201	GTTACCAATG CAATGGTTAC	CTTAATCAGT GAATTAGTCA				
7251	CGTTCATCCA GCAAGTAGGT	TAGTTGCCTG ATCAACGGAC			- · · ·	·
7301	GGAGGGCTTA CCTCCCGAAT	CCATCTGGCC GGTAGACCGG				
7351	GCTCACCGGC CGAGTGGCCG	TCCAGATTTA AGGTCTAAAT				
7401	CTCGCGTCTT	CACCAGGACG	TTGAAATAGG	CGGAGGTAGG	TCAGATAATT	
7451		CTTCGATCTC	ATTCATCAAG	CGGTCAATTA	TCAAACGCGT	
7501	ACGTTGTTGC TGCAACAACG	CATTGCTACA GTAACGATGT	GGCATCGTGG CCGTAGCACC	TGTCACGCTC ACAGTGCGAG	GTCGTTTGGT CAGCAAACCA	
7551	ATGGCTTCAT TACCGAAGTA	TCAGCTCCGG AGTCGAGGCC	TTCCCAACGA AAGGGTTGCT	TCAAGGCGAG AGTTCCGCTC	TTACATGATC AATGTACTAG	

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7	601		TGCAAAAAAG ACGTTTTTTC				
7	651		GTTGGCCGCA CAACCGGCGT				
7	701		TTACTGTCAT AATGACAGTA				
7	751		ACCAAGTCAT TGGTTCAGTA				
7	801	CGAGAACGGG	GGCGTCAATA CCGCAGTTAT	GCCCTATTAT	GGCGCGGTGT	ATCGTCTTGA	
7	851	TTAAAAGTGC AATTTTCACG	TCATCATTGG AGTAGTAACC	AAAACGTTCT TTTTGCAAGA	TCGGGGCGAA AGCCCCGCTT	AACTCTCAAG	
7	901	GATCTTACCG	CTGTTGAGAT GACAACTCTA	CCAGTTCGAT	GTAACCCACT	CGTGCACCCA	
	- [] - '9 <u>51</u> -		AGCATCTTTT TCGTAGAAAA				
8			AAAATGCCGC TTTTACGGCG				
8	3051 		ATACTCTTCC TATGAGAAGG				
. 8	31 0 1 		CATGAGCGGA GTACTCGCCT				
8	3151		TTCCGCGCAC AAGGCGCGTG				

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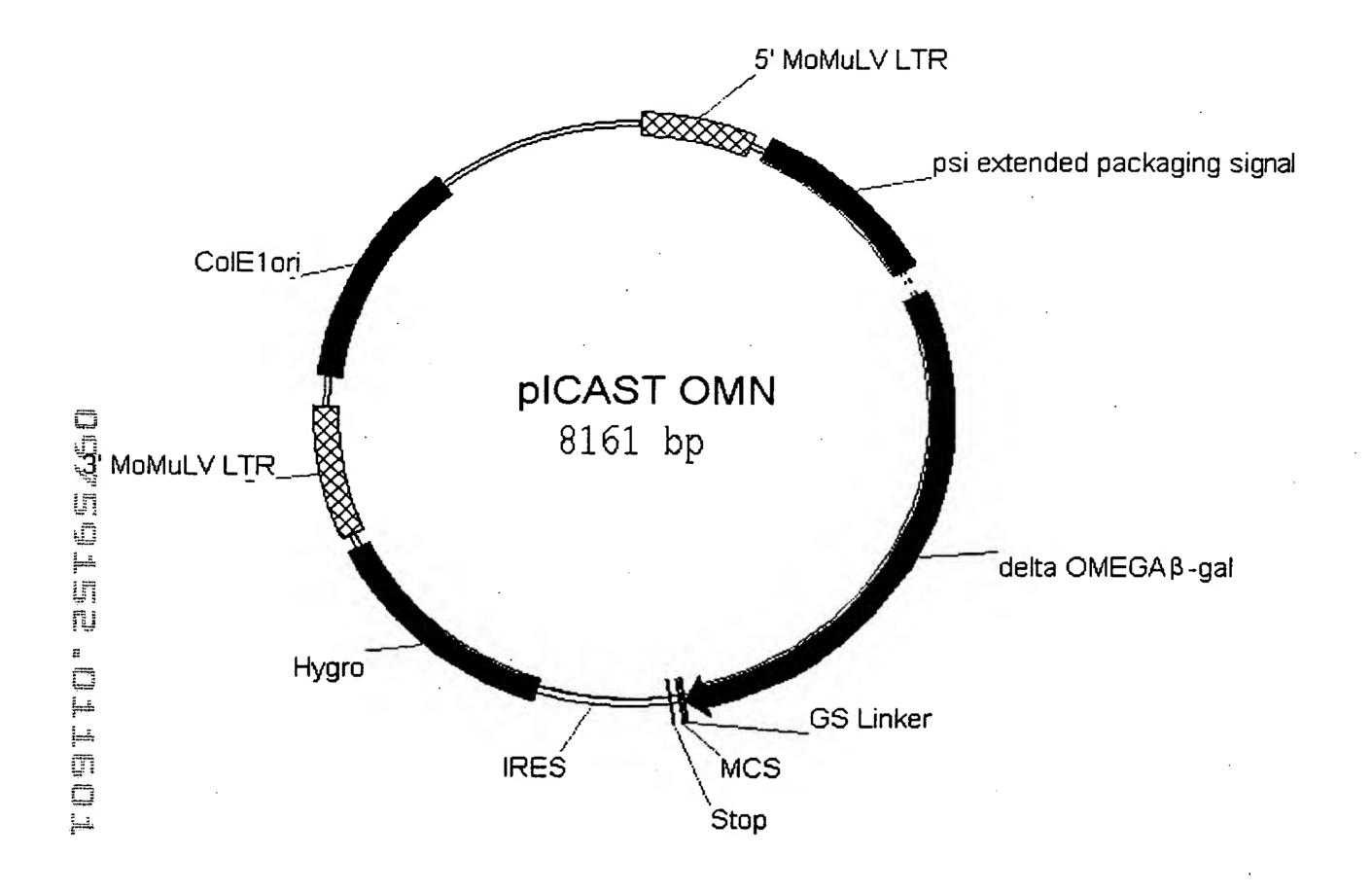


Figure 13A

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101	GGATATCTGT CCTATAGACA				AAGAACAGAT TTCTTGTCTA
151	GGTCCCCAGA CCAGGGGTCT				ACCATCAGAT TGGTAGTCTA
201	GTTTCCAGGG CAAAGGTCCC				TTATTTGAAC AATAAACTTG
251	TAACCAATCA ATTGGTTAGT	CAAGCGAAGA	GCGAAGACAA	GCGCGCGAAG	ACGAGGGCT
301	GCTCAATAAA CGAGTTATTT	TCTCGGGTGT	ACCCCTCACT TGGGGAGTGA	GCCCGCGGT	GTCCTCCGAT CAGGAGGCTA
-67 - 3 <u>51</u> 	TGACTGAGTC ACTGACTCAG	GCCCGGGTAC CGGGCCCATG	CCGTGTATCC GGCACATAGG	AATAAACCCT TTATTTGGGA	CTTGCAGTTG GAACGTCAAC
401	CATCCGACTT	GTGGTCTCGC	TGTTCCTTGG	GAGGGTCTCC	TCTGAGTGAT AGACTCACTA
451					CCGGGATCGG GGCCCTAGCC
5 01	GAGACCCCTG CTCTGGGGAC		TGGCTGGGTG	GTGGCCCTCC	
-	AGCAACTTAT				ACTGATTTTA TGACTAAAAT
601	TGCGCCTGCG ACGCGGACGC				CTGGCGGACC GACCGCCTGG
	CGTGGTGGAA GCACCACCTT				CTGGGAGACG GACCCTCTGC
		AAACCCCCGG	CAAAAACACC	GGGCTGGACT	GGAAGGGAGT CCTTCCCTCA
	CGATGTGGAA GCTACACCTT	TCCGACCCCG AGGCTGGGGC	TCAGGATATG AGTCCTATAC	TGGTTCTGGT ACCAAGACCA	
801	AACCTAAAAC TTGGATTTTG	AGTTCCCGCC TCAAGGGCGG	TCCGTCTGAA AGGCAGACTT	TTTTTGCTTT AAAAACGAAA	CGGTTTGGAA
	CCGAAGCCGC GGCTTCGGCG	GCGTCTTGTC CGCAGAACAG	TGCTGCAGCA ACGACGTCGT	TCGTTCTGTG AGCAAGACAC	TTGTCTCTGT
	CTGACTGTGT GACTGACACA	TTCTGTATTT AAGACATAAA	GTCTGAAAAT CAGACTTTTA	TAGGGCCAGA ATCCCGGTCT	CTGTTACCAC

951	TCCCTTAAGT AGGGAATTCA			AGATGTCGAG TCTACAGCTC		
1001				GTTGGGTTAC CAACCCAATG		
1051	GCAGAATGGC CGTCTTACCG			CCGCGAGACG GGCGCTCTGC		·
1101	CCGAGACCTC GGCTCTGGAG			GGTCTTTTCA CCAGAAAAGT		
1151	ATGGACACCC TACCTGTGGG			TGACCTGGGA ACTGGACCCT		
1201				GTACACCCTA CATGTGGGAT		
1251				CCTTGAACCT GGAACTTGGA		
1301		TAGGAGGGAA	ATAGGTCGGG	AGTGAGGAAG		
1351	GGCCGCTCTA CCGGCGAGAT	GCCCATTAAT CGGGTAATTA	ACGACTCACT TGCTGAGTGA	ATAGGGCGAT TATCCCGCTA	TCGAACACCA AGCTTGTGGT	
=		TCATCATCAC	GTCGACGAAC	AGAAACTCAT	TTCCGAAGAA	
•	GACCTACTCG CTGGATGAGC					
1501	ACGTCGTGAC TGCAGCACTG			CCAACTTAAT GGTTGAATTA		
1551	CACATCCCCC GTGTAGGGGG			GCGAAGAGGC CGCTTCTCCG		
1601	CGCCCTTCCC GCGGGAAGGG			GGCGAATGGC CCGCTTACCG		
1651	GTTTCCGGCA CAAAGGCCGT			CTGGCTGGAG GACCGACCTC		
1701	CTGAGGCCGA GACTCCGGCT			ACTGGCAGAT TGACCGTCTA		
1751	GATGCGCCCA CTACGCGGGT		GCACTGGATA	GGGTAATGCC	AGTTAGGCGG	
1801	GTTTGTTCCC CAAACAAGGG					
1851	TTGATGAAAG AACTACTTTC			CGCGAATTAT GCGCTTAATA		

1901		CGTTTCATCT GCAAAGTAGA	CACCACGTTG			
1951		CGTTTGCCGT GCAAACGGCA				
2001		CCGCCTCGCG GGCGGAGCGC				
2051		ATCAGGATAT TAGTCCTATA				
2101		CATAAACCGA GTATTTGGCT				
2151		TGATGATTTC ACTACTAAAG				
2201		AGTTGCGTGA TCAACGCACT	GATGGATGCC		GAAATACCGT	
22 5]	CCCACTTTGC	CAGGTCGCCA GTCCAGCGGT	GCGGCACCGC CGCCGTGGCG	GCCTTTCGGC CGGAAAGCCG	GGTGAAATTA	
23 91	TCGATGAGCG AGCTACTCGC	TGGTGGTTAT ACCACCAATA	GCCGATCGCG CGGCTAGCGC	TCACACTACG AGTGTGATGC	TCTGAACGTC	
2351	GAAAACCCGA CTTTTGGGCT	AACTGTGGAG TTGACACCTC	CGCCGAAATC GCGGCTTTAG	CCGAATCTCT GGCTTAGAGA	ATCGTGCGGT TAGCACGCCA	
24 01	GGTTGAACTG		ACGGCACGCT	GATTGAAGCA	GAAGCCTGCG	
2451		CCGCGAGGTG GGCGCTCCAC				
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2601		CAACTTTAAC GTTGAAATTG				
2651		ACACGCTGTG TGTGCGACAC				
2701	AGCCAATATT TCGGTTATAA	GAAACCCACG CTTTGGGTGC				
2751		CGATGGCCGC	TACTCGCTTG	CGCATTGCGC	TTACCACGTC	*
2801	GCGCTAGCAT	TAGTGGGCTC	ACACTAGTAG	ACCAGCGACC	CCTTACTTAG	

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2851	TCCGGTGCCG	GCTAATCACG CGATTAGTGC	TGCGCGACAT	AGCGACCTAG	TTTAGACAGC	
2901		CCCGGTGCAG GGGCCACGTC				
2951	ACCGATATTA TGGCTATAAT	TTTGCCCGAT AAACGGGCTA				
3001	CCCGGCTGTG GGGCCGACAC	CCGAAATGGT GGCTTTACCA				
3051	AGACGCGCCC TCTGCGCGGG	GCTGATCCTT CGACTAGGAA				
	CTTGGCGGTT GAACCGCCAA	TCGCTAAATA AGCGATTTAT	GACCGTCCGC	AAAGCAGTCA		
31,51		TTCGTCTGGG AAGCAGACCC				
3201	TACTTTTGCC	CAACCCGTGG GTTGGGCACC	AGCCGAATGC	CGCCACTAAA		
3251 FU		GCCAGTTCTG CGGTCAAGAC				
33 <u>01</u> 	CGGCGTAGGT	GCGCTGACGG CGCGACTGCC	TTCGTTTTGT	GGTCGTCGTC		
3351	TCCGTTTATC	CGGGCAAACC GCCCGTTTGG				
3401	CATAGCGATA GTATCGCTAT	ACGAGCTCCT TGCTCGAGGA				
3451	GCTGGCAAGC CGACCGTTCG	GGTGAAGTGC CCACTTCACG				
3501		GCCTGAACTA CGGACTTGAT				
3551	CTCACAGTAC GAGTGTCATG	GCGTAGTGCA CGCATCACGT				
3601	GCACATCAGC CGTGTAGTCG	GCCTGGCAGC CGGACCGTCG			- · · · -	
3651		GCGCAGGGTG	CGGTAGGGCG	TAGACTGGTG	GTCGCTTTAC	·
3701		AGCTCGACCC	ATTATTCGCA	ACCGTTAAAT	TGGCGGTCAG	
3751		AGTGTCTACA	CCTAACCGCT	ATTTTTTGTT	GACGACTGCG	
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3801		TCAGTTCACC AGTCAAGTGG				
3851	AGGCCTTGGC TCCGGAACCG	GCGCCGGATC CGCGGCCTAG				
3901		ACTGATTAGA TGACTAATCT				
3951	CGGTTATTTT	CCACCATATT GGTGGTATAA				
4001	ACCTGGCCCT TGGACCGGGA	CAGAAGAACT		ATCCCCAGAA		
4051 - = -	CCAAAGGAAT GGTTTCCTTA	GCAAGGTCTG CGTTCCAGAC				
4101		GAAGACAAAC CTTCTGTTTG				
415 <u>1</u>		CCTGGCGACA GGACCGCTGT				
4201		TGCAAAGGCG ACGTTTCCGC				
	ATAGTTGTGG TATCAACACC					
4301,		GCCCAGAAGG CGGGTCTTCC				
4351		ATGCTTTACA TACGAAATGT				·
4401		CCACGGGGAC GGTGCCCCTG				
4451		AAGCCTGAAC TTCGGACTTG			AAGTTTCTGA TTCAAAGACT	
4501		GCTGTCGCAG	AGGCTGGACT	ACGTCGAGAG	CCTCCCGCTT	
4551	CTTAGAGCAC	GAAAGTCGAA	GCTACATCCT	CCCGCACCTA	ATGTCCTGCG TACAGGACGC	
4601	CCATTTATCG	ACGCGGCTAC	CAAAGATGTT	TCTAGCAATA	GTTTATCGGC CAAATAGCCG	
4651	TGAAACGTAG	CCGGCGCGAG	GGCTAAGGCC	TTCACGAACT	CATTGGGGAA GTAACCCCTT	
4701		CGGACTGGAT	AACGTAGAGG	GCGGCACGTG	AGGGTGTCAC TCCCACAGTG	
						· · ·

4751	GTTGCAAGAC CAACGTTCTG			CGCTGTTCTG GCGACAAGAC		
4801	CGGAGGCCAT GCCTCCGGTA			ATCTTAGCCA TAGAATCGGT		
4851	TTCGGCCCAT AAGCCGGGTA			CAATACACTA GTTATGTGAT		
4901	TTTCATATGC AAAGTATACG			GTATCACTGG CATAGTGACC		
4951	TGGACGACAC ACCTGCTGTG			AGGCTCTCGA TCCGAGAGCT		
5001 - 🖵 -	CTTTGGGCCG GAAACCCGGC	TCCTGACGGG		GTGGAGCACG		
50 5 11 - 151 -				CCGCATAACA GGCGTATTGT		
	ACTGGAGCGA TGACCTCGCT	GGCGATGTTC CCGCTACAAG	GGGGATTCCC CCCCTAAGGG	AATACGAGGT TTATGCTCCA	CGCCAACATC GCGGTTGTAG	
5151				GAGCAGCAGA CTCGTCGTCT	CGCGCTACTT GCGCGATGAA	
52 01 - 				GCCGCGGCTC CGGCGCCGAG	CGGGCGTATA GCCCGCATAT	
52 51 - 					TGACGGCAAT ACTGCCGTTA	
5301 	AAGCTACTAC	GTCGAACCCG	CGTCCCAGCT	ACGCTGCGTT	TCGTCCGATC AGCAGGCTAG	
5351	GCCTCGGCCC	TGACAGCCCG	CATGTGTTTA	GCGGGCGTCT	AGCGCGGCCG TCGCGCCGGC	
5401	AGACCTGGCT	ACCGACACAT	CTTCATGAGC	GGCTATCACC	AAACCGACGC TTTGGCTGCG	
5451	GGGTCGTGAG	CAGGCTCCCG	TTTCCTTATC	TCATCTACGG	GACCGGGATC	
5501	ATAGCTATTT	TATTTTCTAA	AATAAATCAG	AGGTCTTTTT	GGGGGGAATG	
5551	TTTCTGGGGT	GGACATCCAA	ACCGTTCGAT	CGAATTCATT		
5601	GTTCCGTACC	TTTTTATGTA	TTGACTCTTA	TCTCTTCAAG		
5651					ATCTGTGGTA A TAGACACCAT	· ·

5701	AGCAGTTCCT TCGTCAAGGA			ACAGATGGAA TGTCTACCTT		
5751	TGGGCCAAAC ACCCGGTTTG			TTCCTGCCCC AAGGACGGGG	CCGAGTCCCG	
5801	CAAGAACAGA GTTCTTGTCT			GCCCTCAGCA CGGGAGTCGT		
5851	AACCATCAGA TTGGTAGTCT			GACCTGAAAT CTGGACTTTA		
5901	CTTATTTGAA GAATAAACTT			TCGCTTCTGT AGCGAAGACA		
5951 	CTGCTCCCCG GACGAGGGGC			AACCCCTCAC TTGGGGAGTG		
6001				CCCGTGTATC GGGCACATAG		
6051 -				CTGTTCCTTG GACAAGGAAC		
61 0 1				GTCTTTCATT CAGAAAGTAA		
61 <u>51</u> - <u> -1</u>				AGTATTTACA TCATAAATGT		
62 <u>0</u> 1				GGGGAGAGGC CCCCTCTCCG		
6251		_		CTCGCTGCGC GAGCGACGCG		
6301				AGGCGGTAAT TCCGCCATTA		
6351				ATGTGAGCAA TACACTCGTT		
6401	AAAGGCCAGG TTTCCGGTCC			GCTGGCGTTT CGACCGCAAA		
6451				GACGCTCAAG CTGCGAGTTC	TCAGAGGTGG AGTCTCCACC	
6501		GTCCTGATAT	TTCTATGGTC	CGCAAAGGGG	GACCTTCGAG	
6551			•		TACCTGTCCG ATGGACAGGC	
6601					ACGCTGTAGG TGCGACATCC	

6651		CGGTGTAGGT GCCACATCCA				
6701		CAGCCCGACC GTCGGGCTGG				
6751	AGTCCAACCC TCAGGTTGGG	GGTAAGACAC CCATTCTGTG				
6801	AACAGGATTA TTGTCCTAAT	GCAGAGCGAG CGTCTCGCTC				
6851		AACTACGGCT TTGATGCCGA				
6901 	GAGACGACTT	CGGTCAATGG	AAGCCTTTTT	CTCAACCATC	GAGAACTAGG	
6951		CCACCGCTGG				
	CCGTTTGTTT	GGTGGCGACC	ATCGCCACCA	AAAAAACAAA	CGTTCGTCGT	
70 61 	CTAATGCGCG	AGAAAAAAAG TCTTTTTTC	CTAGAGTTCT	TCTAGGAAAC		
7051	CGGGGTCTGA					
N.	GCCCCAGACT				CTAAAACCAG	
7101	ATGAGATTAT TACTCTAATA	CAAAAAGGAT GTTTTTCCTA	GAAGTGGATC	TAGGAAAACG	CCGGCGTTTA	
7151	CAATCTAAAG	TATATATGAG	TAAACTTGGT	CTGACAGTTA	CCAATGCTTA	
		ATATATACTC				
7201		CACCTATCTC				
7251	TGCCTGACTC	CCCGTCGTGT	AGATAACTAC	GATACGGGAG	GGCTTACCAT	
	ACGGACTGAG	GGGCAGCACA	TCTATTGATG	CTATGCCCTC	CCGAATGGTA	
7301	CTGGCCCCAG GACCGGGGTC	TGCTGCAATG ACGACGTTAC				
7351	GATTTATCAG	CAATAAACCA	GCCAGCCGGA	AGGGCCGAGC	GCAGAAGTGG	
	CTAAATAGTC	GTTATTTGGT	CGGTCGGCCT	TCCCGGCTCG	CGTCTTCACC	
7401		TTATCCGCCT AATAGGCGGA				
7451	CTAGAGTAAG GATCTCATTC	ATCAAGCGGT	CAATTATCAA	ACGCGTTGCA	ACAACGGTAA	
7501	GCTACAGGCA CGATGTCCGT	TCGTGGTGTC AGCACCACAG	ACGCTCGTCG TGCGAGCAGC	TTTGGTATGG AAACCATACC	CTTCATTCAG GAAGTAAGTC	
7551	CTCCGGTTCC GAGGCCAAGG	CAACGATCAA GTTGCTAGTT	GGCGAGTTAC CCGCTCAATG	ATGATCCCCC TACTAGGGGG	ATGTTGTGCA TACAACACGT	
				. – – – – – – –		

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7601.	AAAAAGCGGT TTTTTCGCCA	TAGCTCCTTC ATCGAGGAAG				
7651	GCCGCAGTGT CGGCGTCACA	TATCACTCAT ATAGTGAGTA				
7701		TCCGTAAGAT AGGCATTCTA				
7751	AGTCATTCTG TCAGTAAGAC	AGAATAGTGT TCTTATCACA				
7801	TCAATACGGG AGTTATGCCC	ATAATACCGC TATTATGGCG				
7851	CATTGGAAAA GTAACCTTTT	CGTTCTTCGG GCAAGAAGCC				
7901		TTCGATGTAA AAGCTACATT				
79 51 		TCACCAGCGT AGTGGTCGCA				
80 <u>01</u>		AAGGGAATAA TTCCCTTATT				
8051		TCAATATTAT AGTTATAATA				
8101	TCGCCTATGT	TATTTGAATG ATAAACTTAC	ATAAATCTTT	TTATTTGTTT		
815 1	GCGCACATTT CGCGTGTAAA	С			 	

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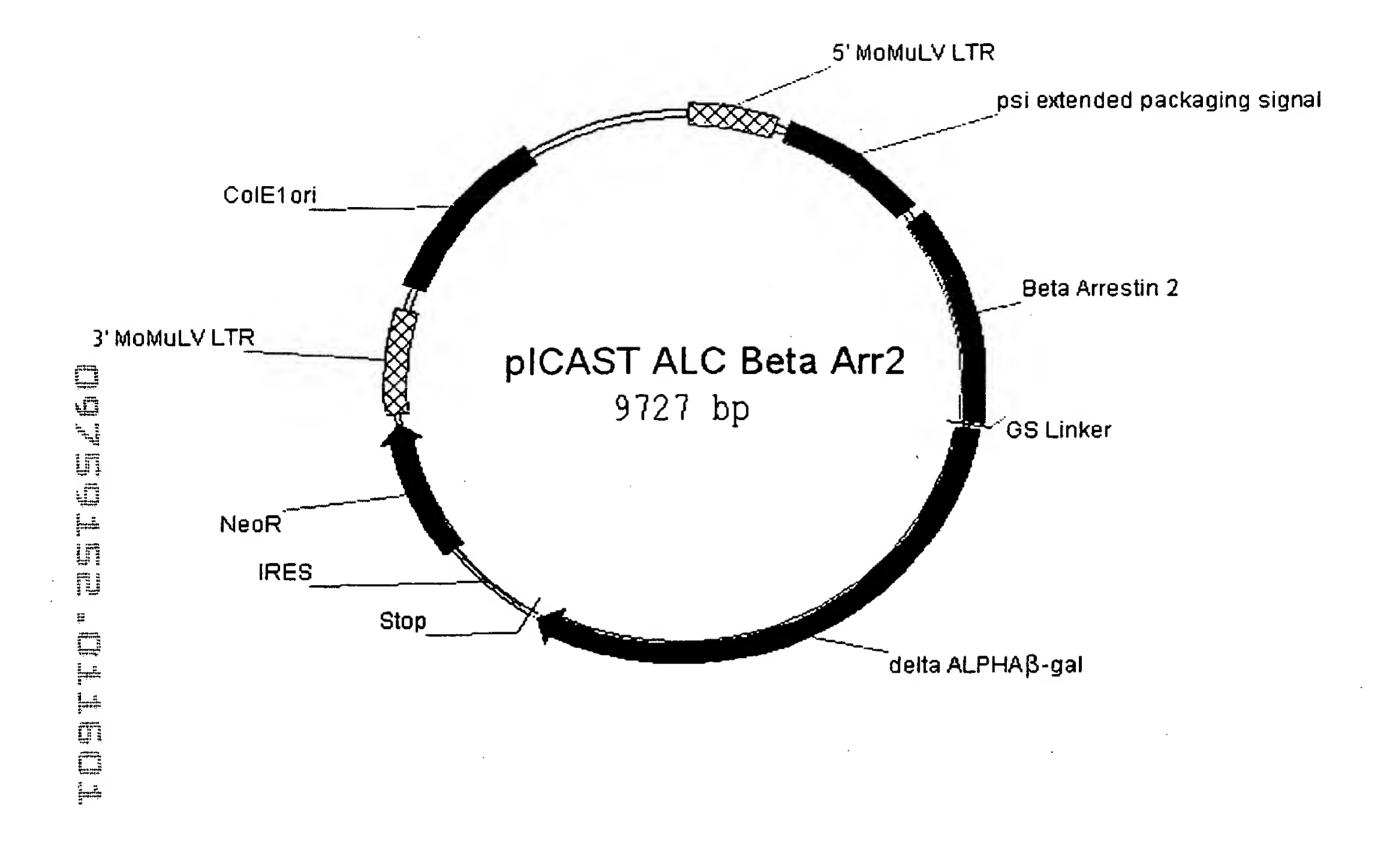


Figure 14

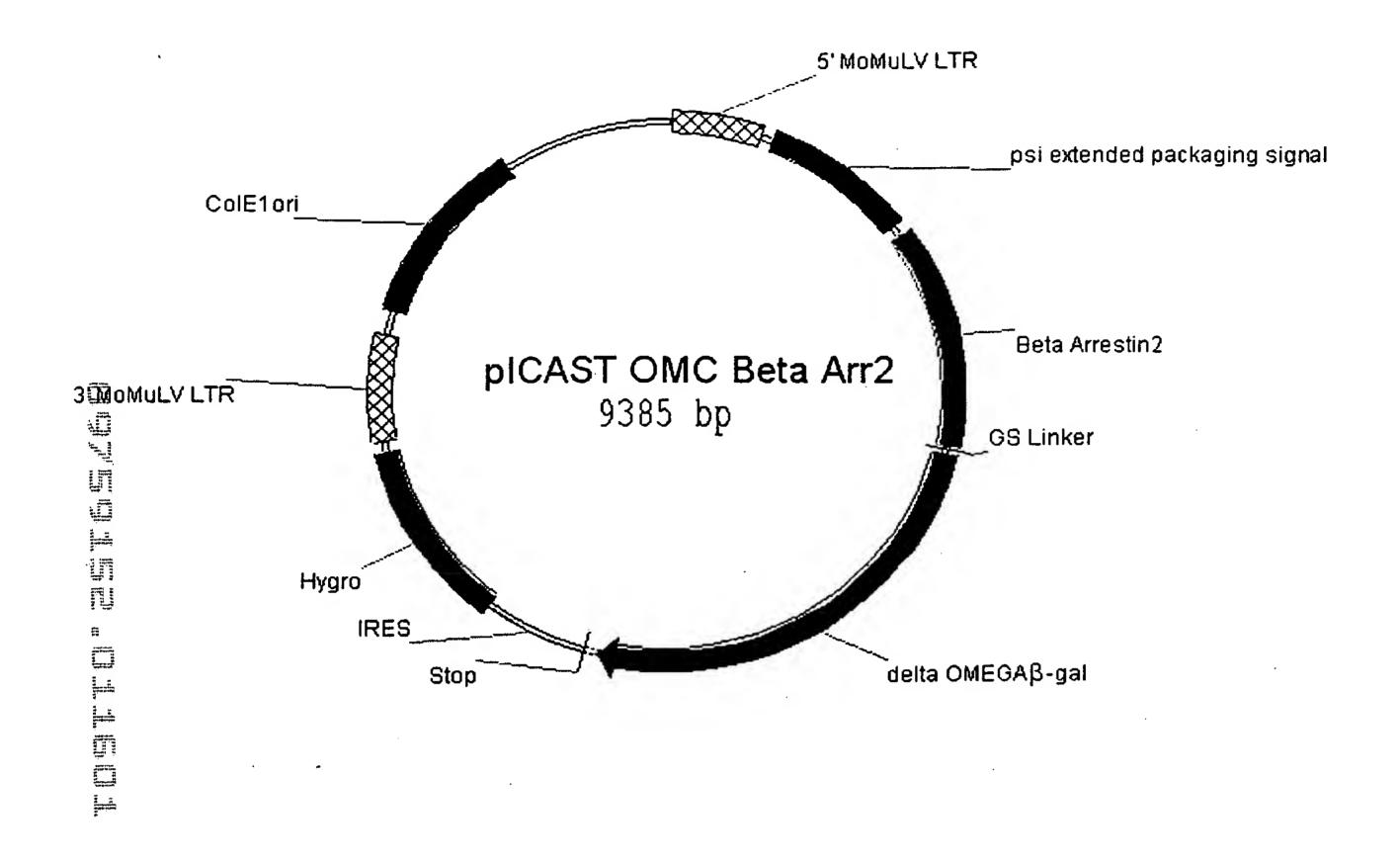


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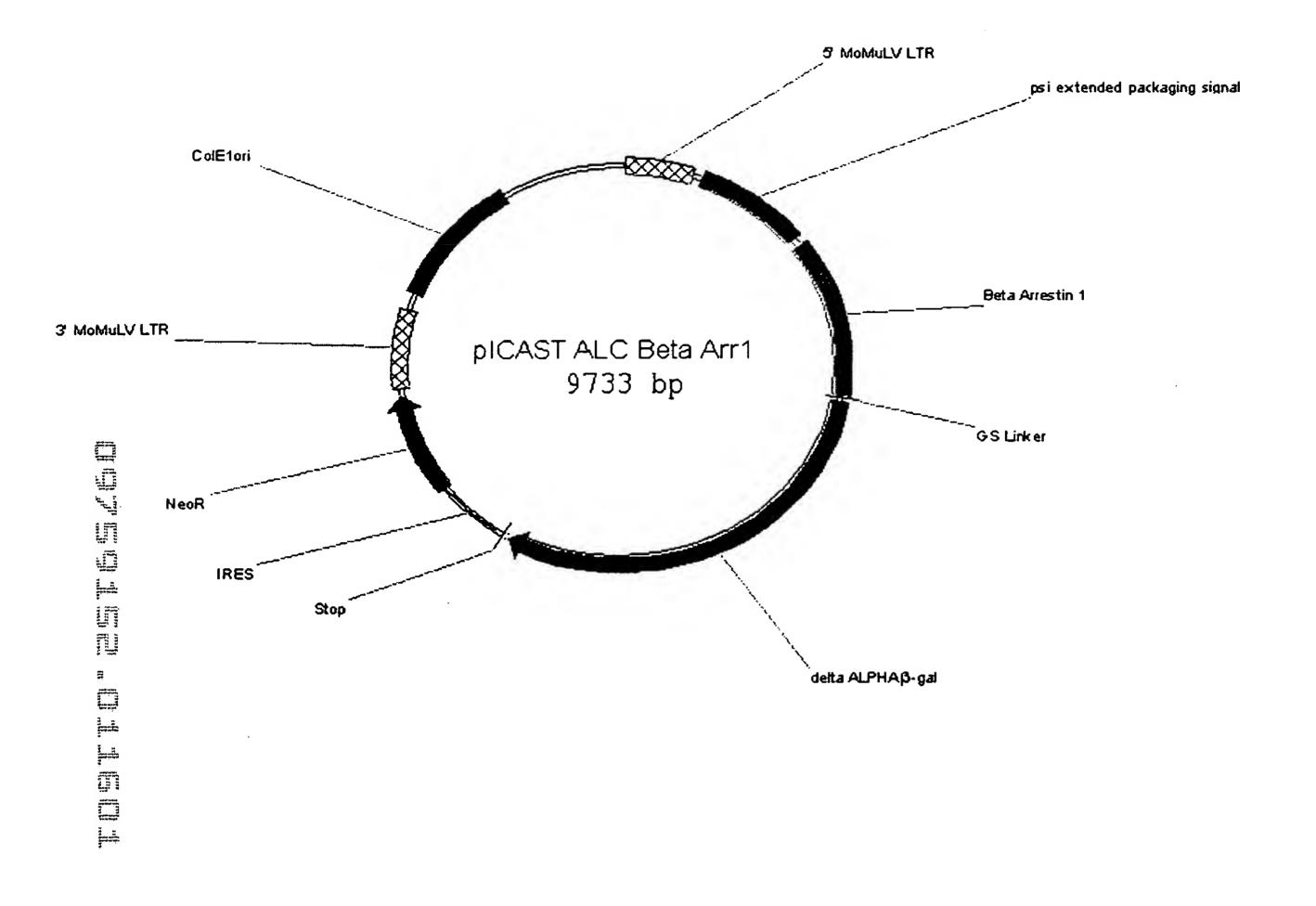


Figure 16

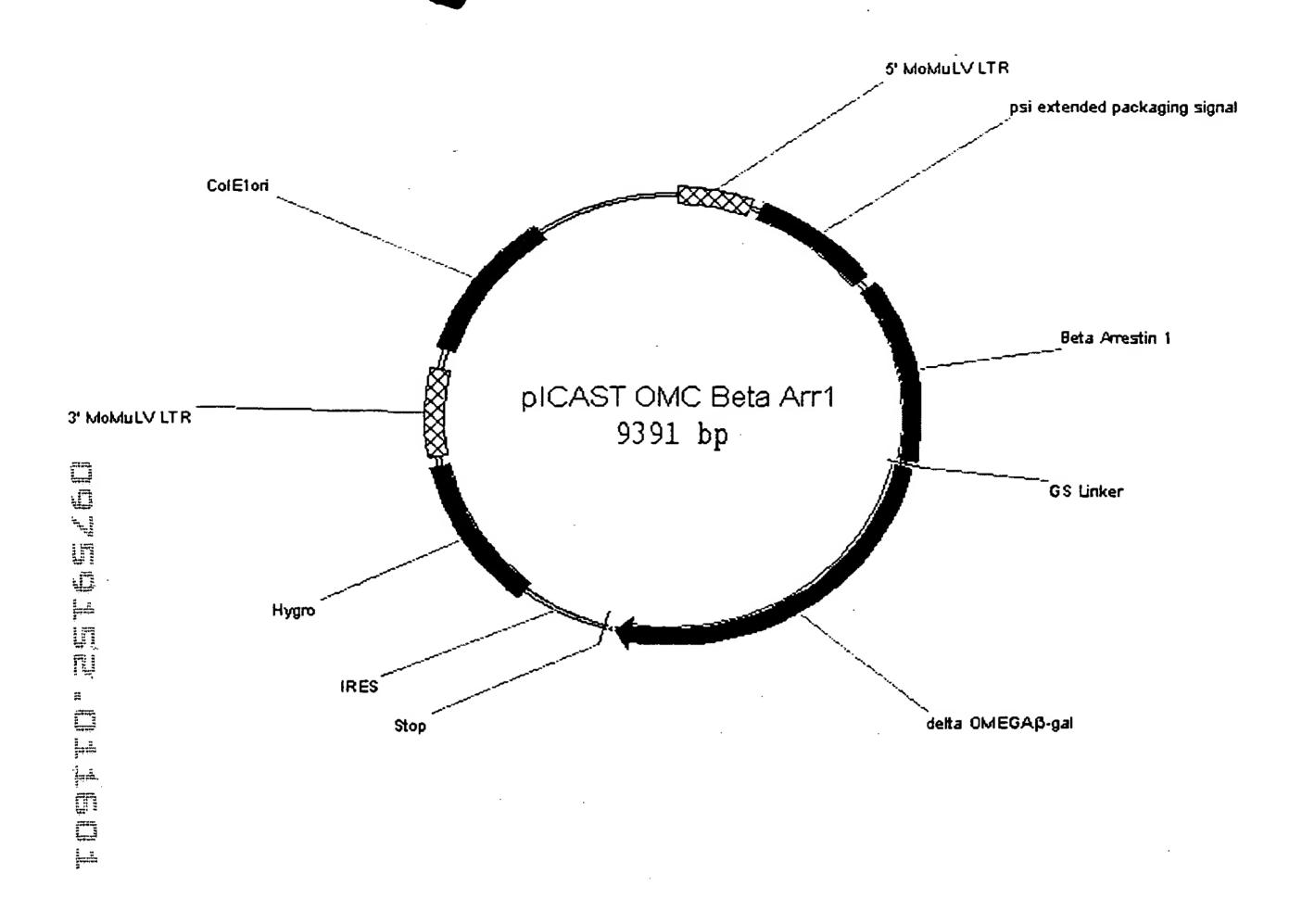


Figure 17

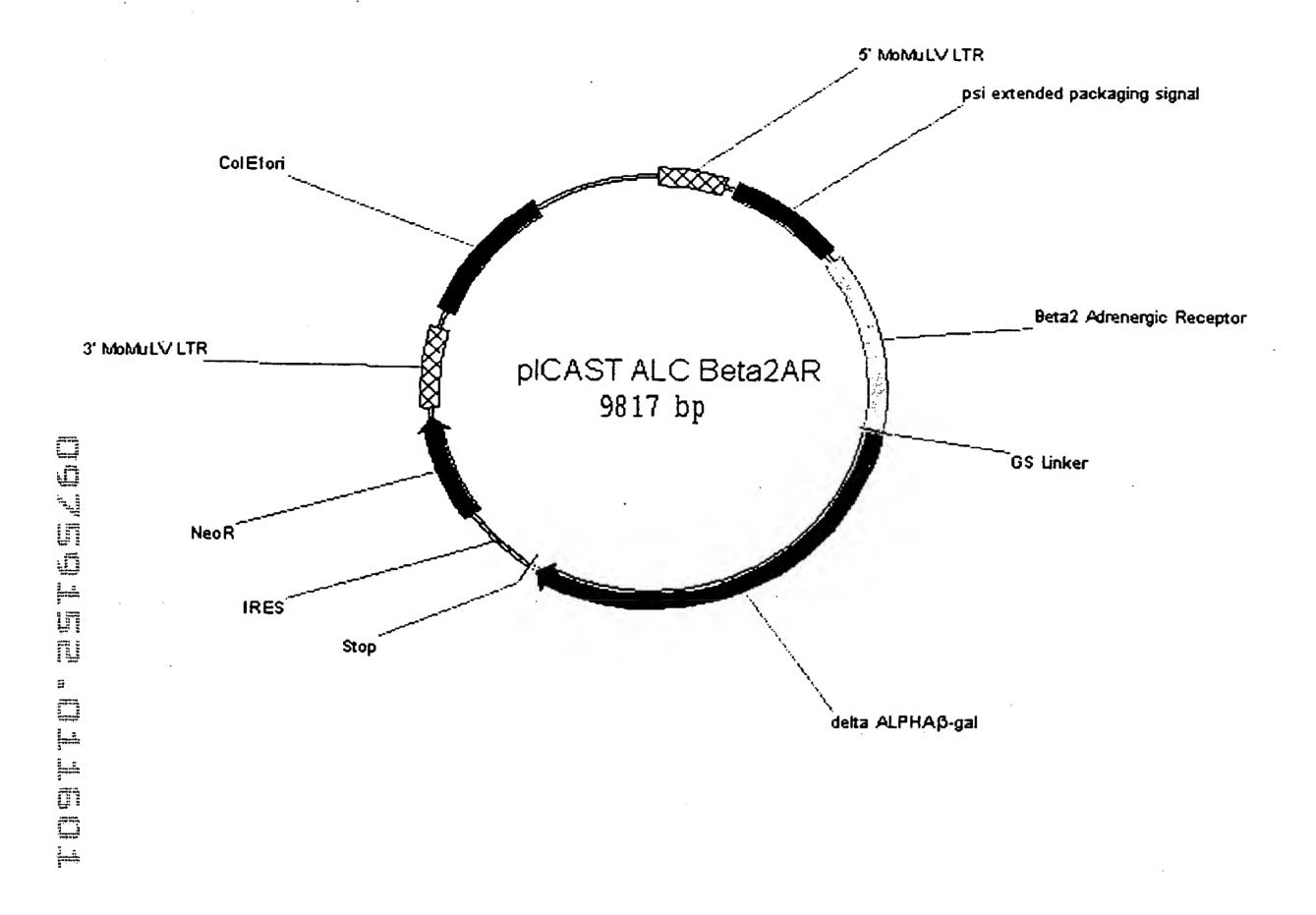


Figure 18

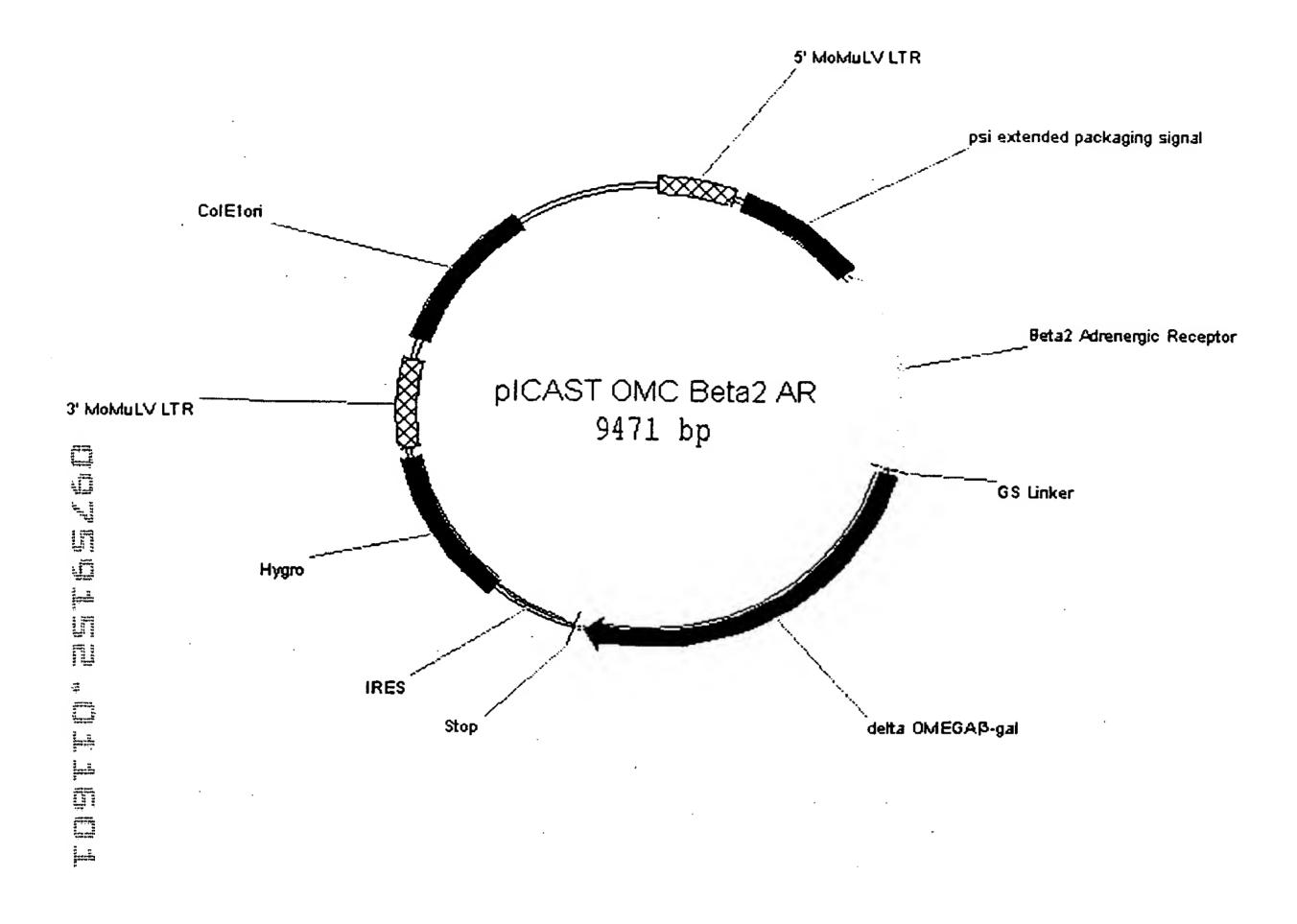


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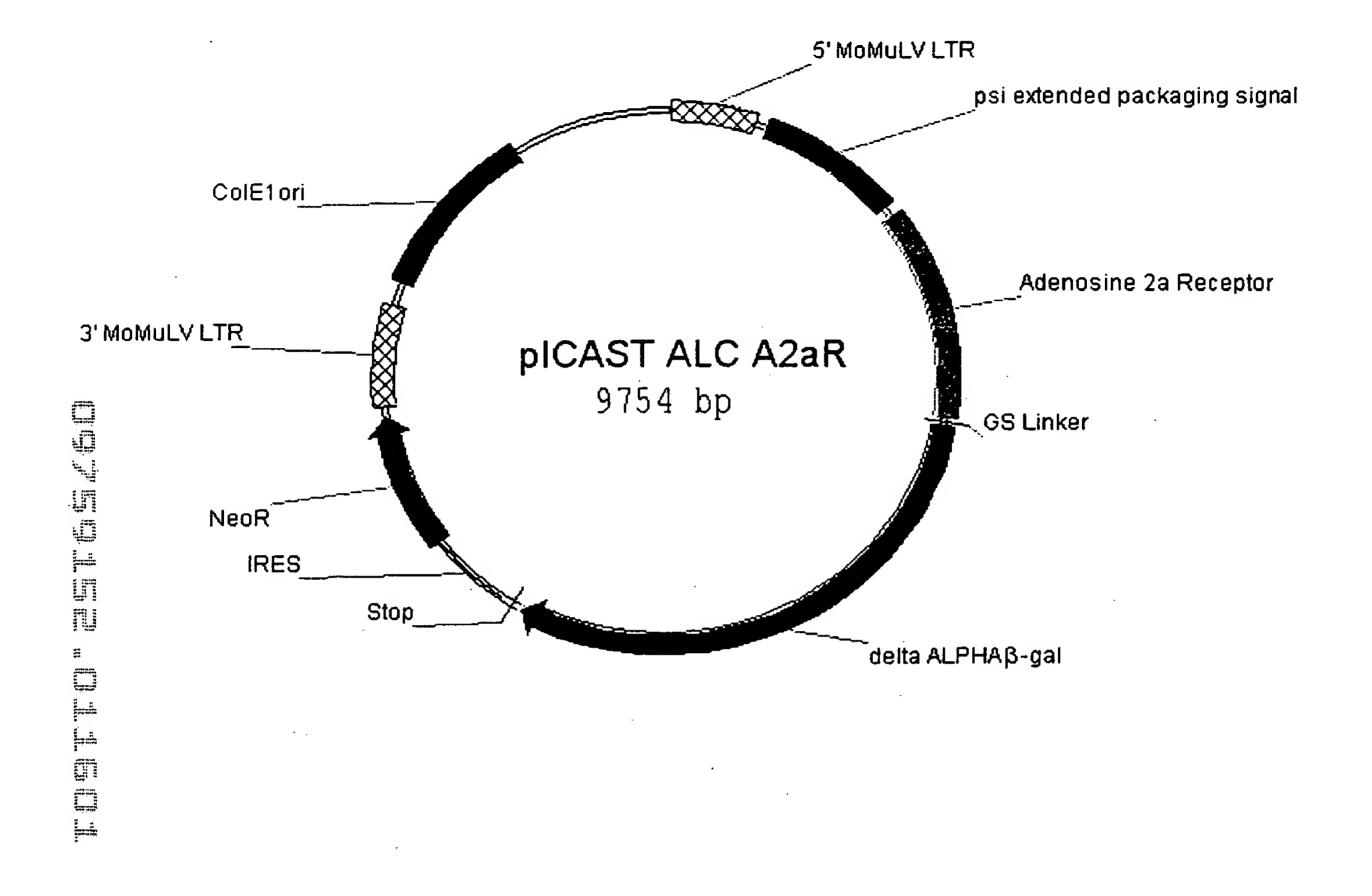


Figure 20

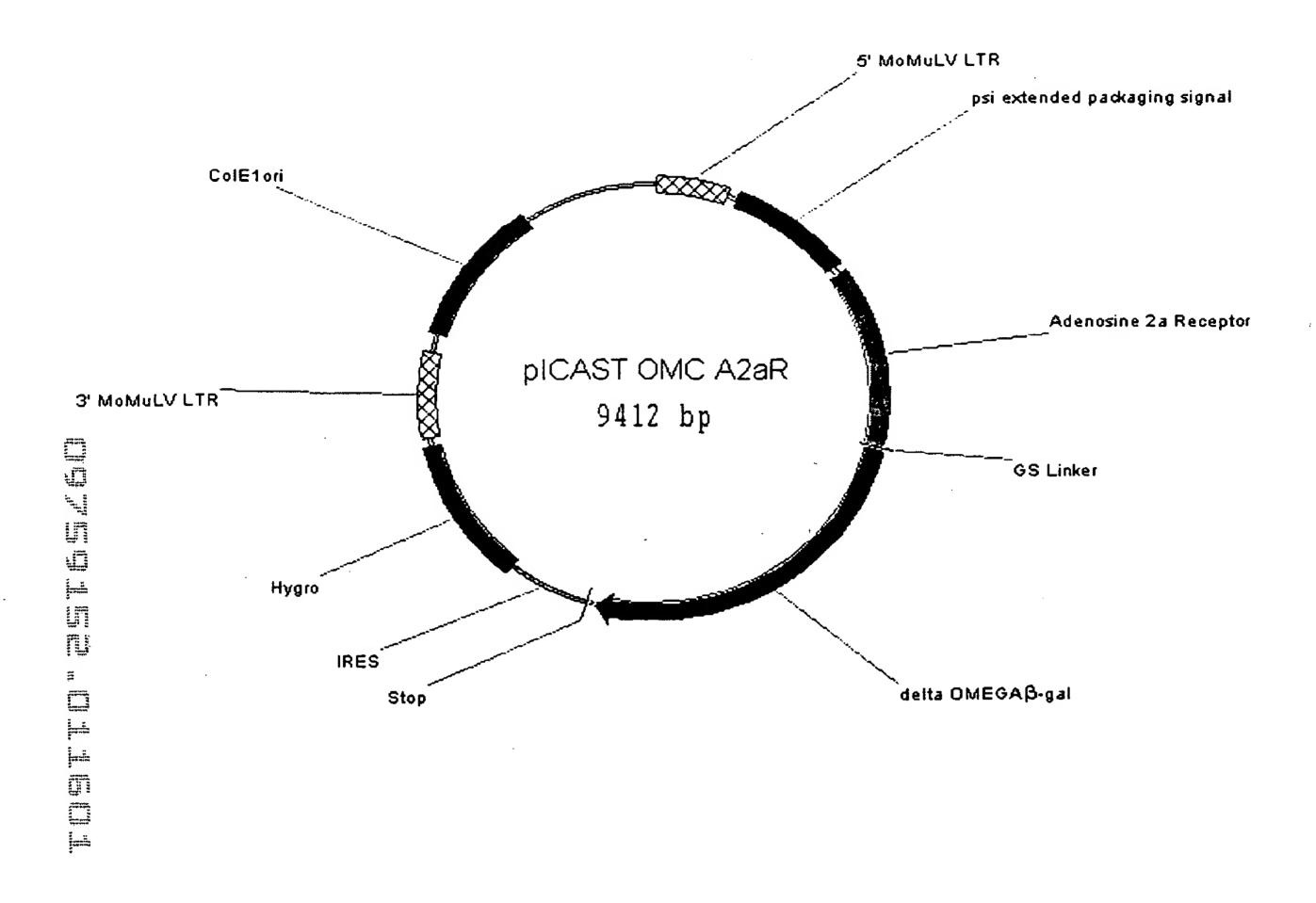


Figure 21

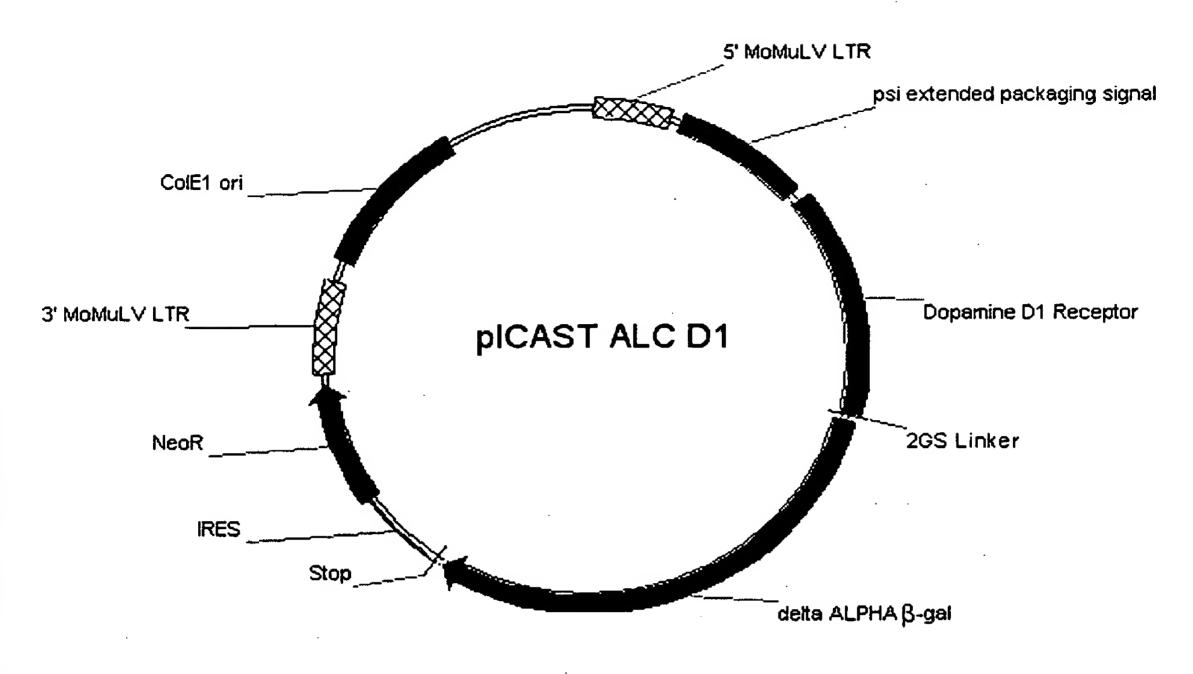


Figure 22

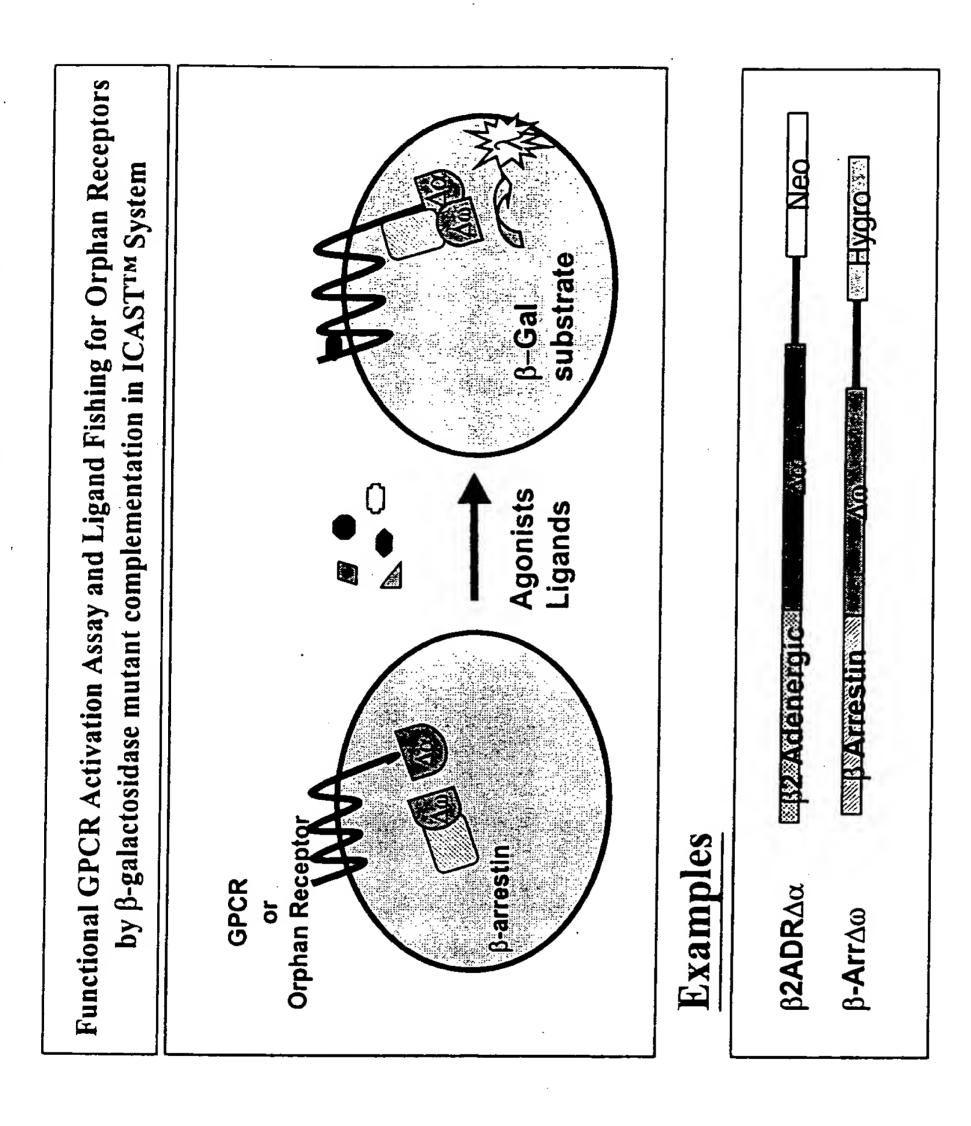
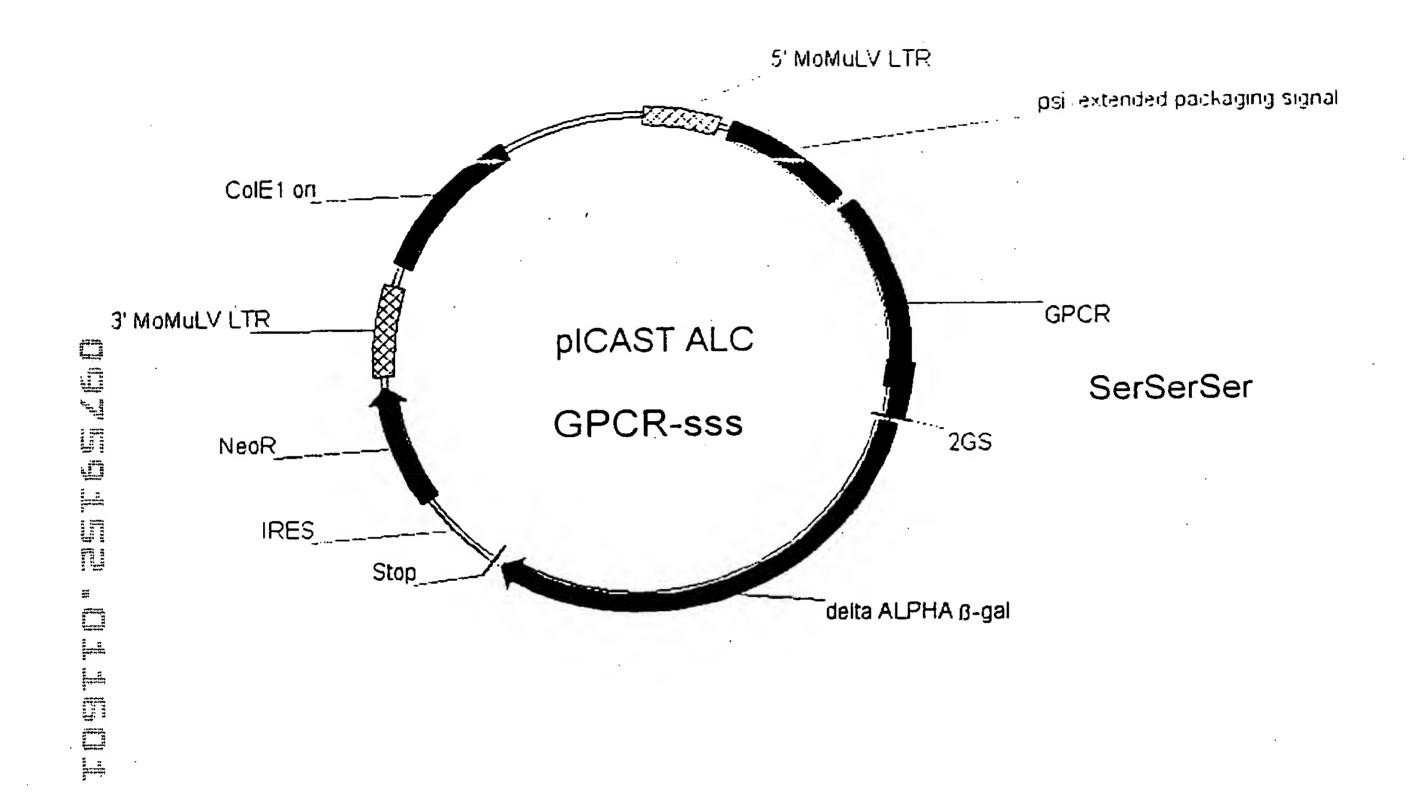
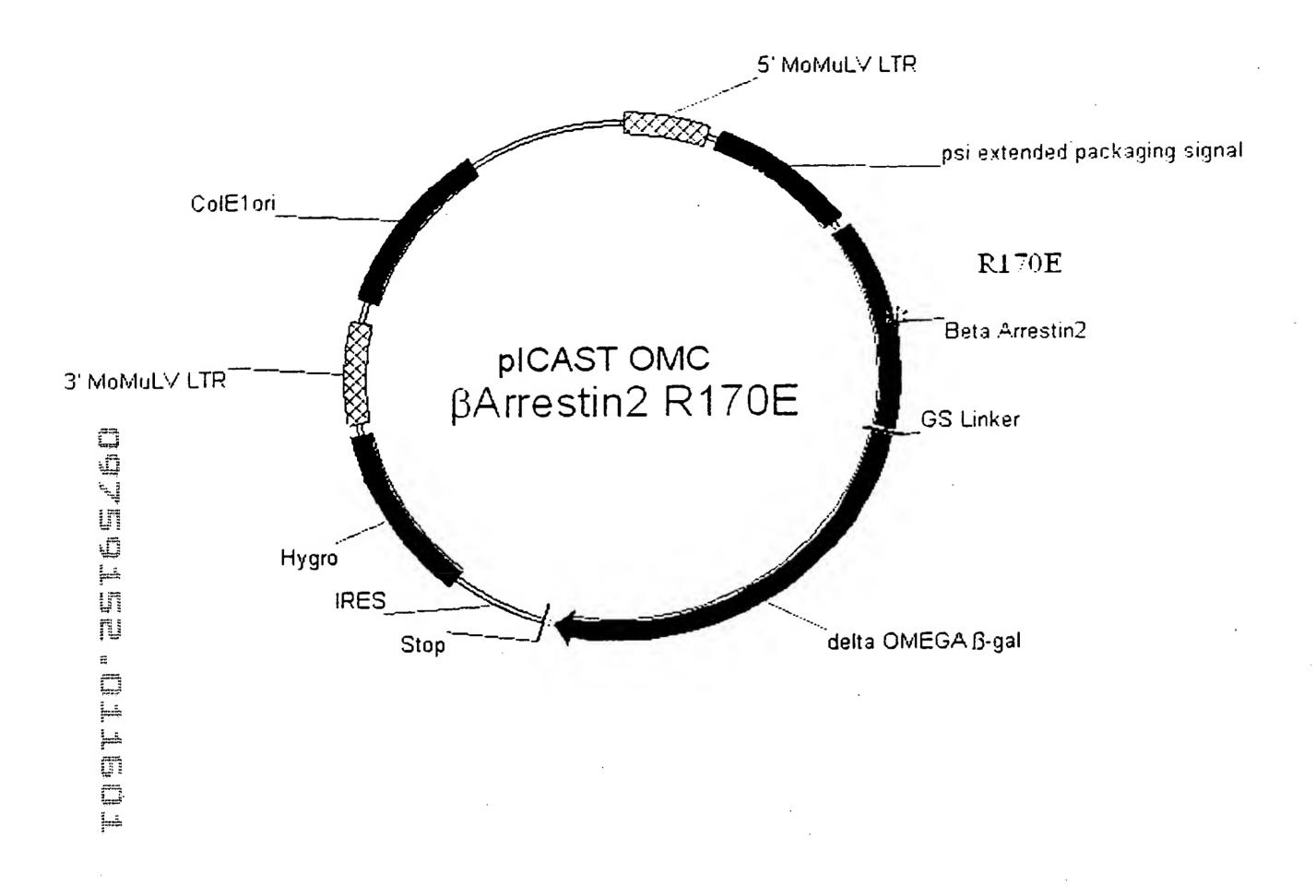


Figure 23



Vector for Expression of a GPCR with inserted Seronine/Threonine amino acid sequences as a fusion with β -gal $\Delta\alpha$.



Vector for Expression of mutant (R170E) $\beta\text{-arrestin2}\;$ as a fusion with $\beta\text{-gal}\;\Delta\omega.$

FIGURE 25

Phosphorylation Insensitive Mutant R170E $\beta\text{-Arrestin2}\Delta\omega$ Binds to $\beta2$ ARA α in Response to Agonist Activation

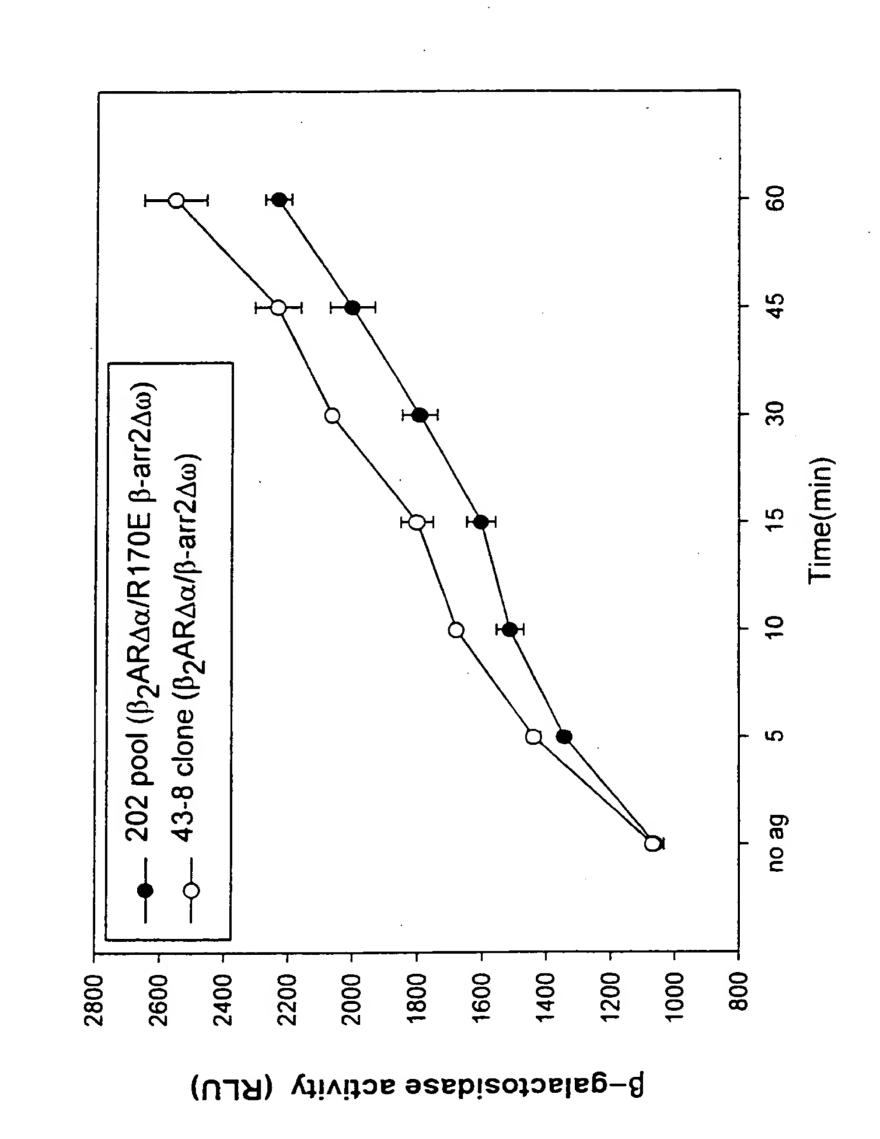
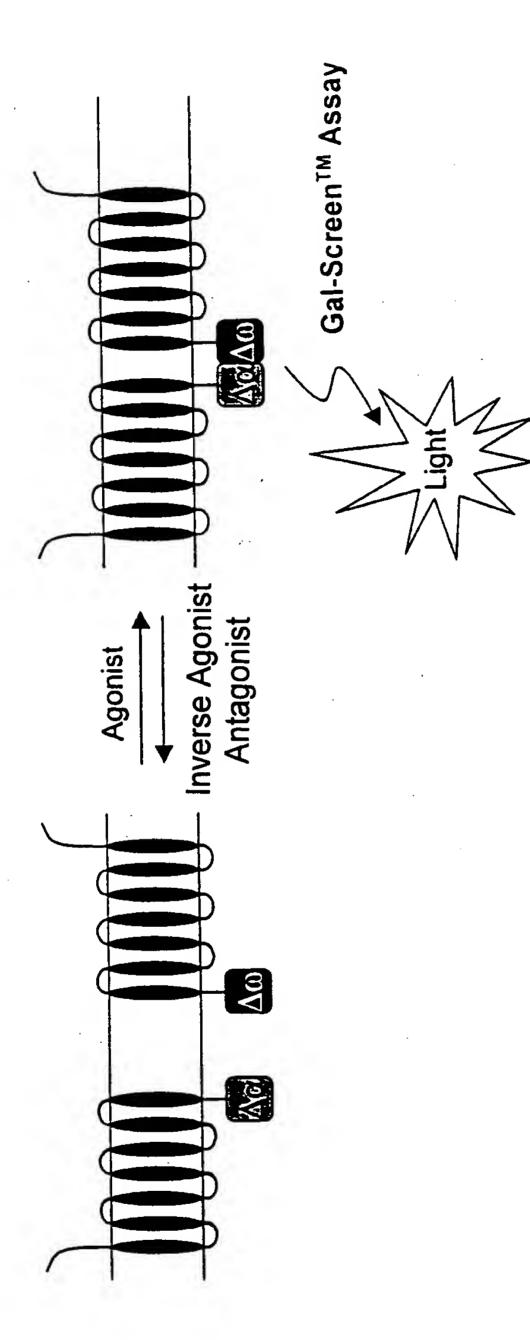
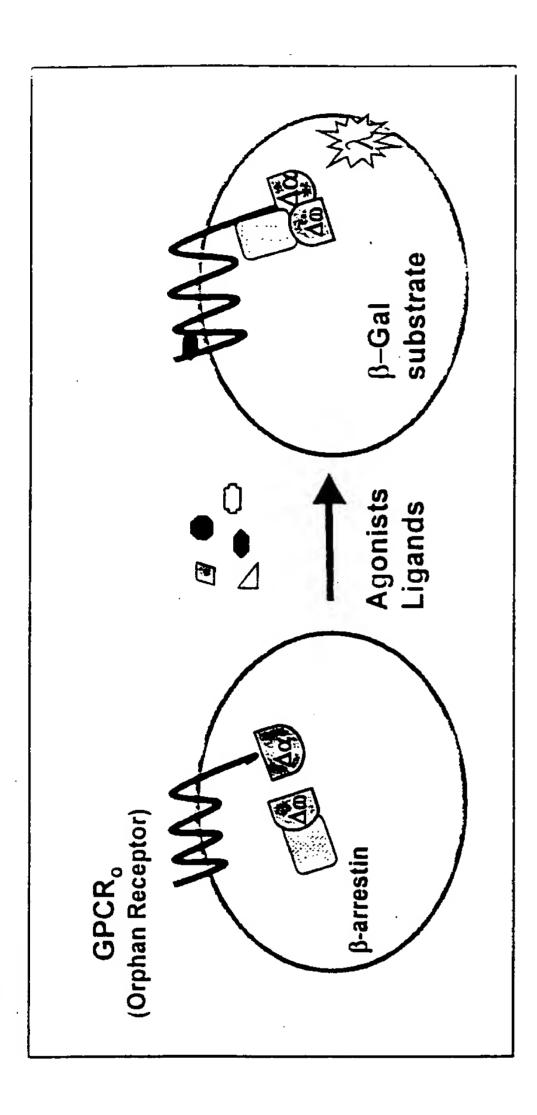


FIGURE 26



GPCR dimerization measured by β -gal complementation

FIGURE 27



GPCRoΔα	Neo Neo
β-ΑιτΔω	RArrestin Instant Axion Man Hygro

Ligand Fishing for Orphan Receptors by β -galactosidase mutant complementation in ICAST^{rM} System

FIGURE 28